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CORRECTED VERSION

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
18 July 2002 (18.07.2002)

PCT

(10) International Publication Number
WO 02/055188 A2

(51) International Patent Classification⁷: **B01J 19/00**,
F16K 3/04

(21) International Application Number: PCT/US02/00370

(22) International Filing Date: 9 January 2002 (09.01.2002)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
60/260,726 10 January 2001 (10.01.2001) US

(63) Related by continuation (CON) or continuation-in-part
(CIP) to earlier application:
US 60/260,726 (CON)
Filed on 10 January 2001 (10.01.2001)

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(81) Designated States (*national*): AE, AG, AL, AM, AT, AU,
AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU,
CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,
LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW,
MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG,
SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ,
VN, YU, ZA, ZM, ZW.

(84) Designated States (*regional*): ARIPO patent (GH, GM,
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW),
Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),
European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR,
GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent
(BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR,
NE, SN, TD, TG).

Published:

— without international search report and to be republished
upon receipt of that report

(48) Date of publication of this corrected version:
31 October 2002

(15) Information about Correction:
see PCT Gazette No. 44/2002 of 31 October 2002, Sec-
tion II

*For two-letter codes and other abbreviations, refer to the "Guid-
ance Notes on Codes and Abbreviations" appearing at the begin-
ning of each regular issue of the PCT Gazette.*

(54) Title: SYSTEM AND PROCESS FOR MICROFLUIDICS-BASED AUTOMATED CHEMISTRY

(57) Abstract: System and process for performing complex reactions such as protein sequencing using small volumes and amounts of reagents, the system in part comprising the use of rotary type selector-switching valve combinations.



WO 02/055188 A2

SYSTEM AND PROCESS FOR MICROFLUIDICS-BASED AUTOMATED CHEMISTRY

The research leading to the present invention was supported, at least in part, by NSF grant DBI-9420123 and Development Funds from NCI Grant P30 CA08748. Accordingly, the Government may have certain rights in the invention.

Background

This invention relates to the field of microfluidics-based automated chemistry.

Many analytical chemical problems involve the analysis of small amounts of a sample. The ability to do chemical reactions in small volumes of liquid is important in such cases as the reaction rate can decrease to commercially unacceptable levels if the concentration of the sample is too low. Additionally, for analytical devices used to monitor the results of such reactions, it is frequently important that the reaction product be in a small volume in order that a detectable concentration be present. The present invention offers a solution to the problem of doing chemical reactions in small volumes, to doing a succession of such reactions, and to doing them on an automated basis.

Although the present invention will be seen to have general chemical applicability, its application to biomedical research is of particular interest. Post-genome research will likely focus on understanding the cellular chemistry, circuitry and communications underlying life's vital processes. Biomedical scientists, for instance, will aim to identify how genetic determinants of disease alter cellular physiology and response to agonists. Predictably, all

this will involve biochemical analysis of larger numbers of samples containing ever lower concentrations of analyte. In most cases, analyses entail multistep procedures, including chemical and enzymatic reactions. Not only will automation prove essential in such cases, it must also be done in fully integrated instruments that incorporate the smallest possible reaction vessels and wetted surfaces. Progress towards this goal has come from nano-fabricated devices¹⁻⁴ ("lab-on-a-chip"), but severe limitations on sample volume may restrict the technology to fast, parallel analysis of abundant and/or amplifiable (e.g. by PCR) molecules. What's needed is automation whereby trace analytes are processed in their entirety. We wanted to construct such a device, initially using chemical protein sequencing as a model system. Such chemistries have been replaced during recent years by mass spectrometry as a means to protein identification.⁵⁻⁸ Yet, the fact remains that chemical analysis can yield rather long stretches of easy to interpret sequence, including on intact proteins, with the caveat that current instruments are at least ten to twenty times less sensitive than most mass analyzers.

Traditionally, protein chemical sequencing is done by stepwise removal of amino acids from the N-terminal end, one at a time. In this method, the *Edman degradation*,^{9,10} phenyl isothiocyanate (PITC) is coupled to the alpha-amino group of the polypeptide to form a phenyl thiocarbamyl (PTC)-derivative; anhydrous acid causes selective release of the PTC-amino, leaving a truncated peptide chain. The resulting anilino thiazolinone amino acid (ATZ-aa) is converted with aqueous acid to a more stable phenyl thiohydantoin amino acid (PTH-aa) and identified. The latter step cannot be done in the presence of polypeptide as it would cause hydrolysis; thus, the ATZ-aa must be extracted. The procedure was partially automated, first by Edman in the 'spinning cup' version,¹¹ and later in Laursen's solid-phase sequencer.¹² Conversion was initially done outside the machine, but later incorporated into the automated process by Wittman-Liebold.¹³ Until then, thin layer or gas chromatography

were used for PTH-aa identification; Hunkapiller and Hood were among the first to routinely use reversed-phase high performance liquid chromatography (RP-HPLC) for this purpose.¹⁴ Around 1980, the gas-phase (GP) sequencer was developed by Hewick and coworkers.¹⁵ Here, wash-out of the peptide from the reaction vessel was prevented by non-covalent immobilization on a glass-fibre disc and by delivering polar liquids as vapors. The process has been further automated by coupling HPLC identification of PTH-aa's 'on-line' with the sequencer; contents of the conversion flask are hereby directly transferred into the LC injector loop. Since then, progress has come from incremental improvements such as femtomole level HPLC detection,¹⁶ rigorous instrument optimization and maintenance routines,¹⁷ and the use of a smaller reaction cartridges.¹⁷⁻¹⁹ Combined with improved micro-preparation of polypeptides,²⁰⁻²² chemical sequencing at the 1-3 picomole level is currently possible and extended sequencing runs with femtomole level signal have been reported.^{16,23,24} Additional modifications to the process have since been suggested that should allow true femtomole level sequencing. All relate to increased sensitivities of amino acid derivative detection. This can be accomplished by miniaturizing the HPLC-based PTH-aa detection system, or by producing modified Edman end-products of higher detect ability, or both.

Whereas femtomole level PTH-aa separations on microbore columns can be done,^{19,25} the use of capillary columns (300-micron ID) is more problematic, as injection volumes of over 0.5 μ L cause baseline disturbances.²⁶ Complete injection (or even 10%) of the sample in a 0.5- μ L volume is not possible from any commercial automated sequencer. This also applies to capillary zone electrophoresis (CZE)-based amino acid-derivative detection systems.²⁷

Considerable effort has been expended to generate fluorescent amino acid derivatives as end-products of Edman chemistry. Fluorescent sequencing is especially appealing since the introduction of sub-attomole amino acid analysis by CZE with laser-induced fluorescence.^{27,28} Yet, again, loading volumes are in the nanoliter range, several orders of

magnitude less than what is typically present in a sequencer flask. Another approach to 'high-detect ability' is the generation of quaternary or tertiary amine thiohydantoin amino acid derivatives, analyzable by mass spectrometry at the low femtomole level.^{29,30} However, both methods never progressed beyond the R&D stage.

Development of any Edman based, femtomole-level technique requires satisfying two major criteria: (i) quantitative transfer, in the smallest possible volume, of amino acid derivatives to the site of analysis; and (ii) reducing chemical background that will impede any ultra-sensitive analytical technique. This can only be accomplished by further miniaturization of the chemistry.

We describe a microfluidics-based instrument, consisting of multiple rotary valves, capillary tubing and miniaturized reaction vessels, for the purpose of performing automated chemical and biochemical reactions on a very small scale. Close to 100% of the reaction end-products are available in a minimal volume ($\leq 5 \mu\text{L}$) inside a pressurized micro-vial for subsequent analysis. This makes the system compatible with capillary HPLC and, in principle, with continuous flow nano-electrospray mass spectrometry. Total control of flow path combinations and directions, temperatures and gas pressures enables precise execution of complex biochemical laboratory procedures. Instrument performance was convincingly demonstrated by partially sequencing 100 femtomoles of an intact protein using classical Edman chemistry in combination with capillary-bore liquid chromatography. To our knowledge, this is the smallest amount of protein ever reported to be successfully analyzed in this way.

Summary of the Invention

The invention is a chemical system and related processes that utilize small-volume rotary selector valves and small-volume rotary switching valves in combination under the control of a computer. The small rotary valves are particular well-suited to computer control. As a result, a single system can be programmed to do a variety of tasks by changing the program but leaving the system's components substantially intact.

As a result, in a general aspect the invention is a system for carrying out one or more chemical reactions, said system comprising a rotary selector valve and a rotary switching valve, each valve under the control of a computer, the internal volumes of the selector and switching valves each being 1.5 μ l or less.

Specific aspects of the invention that are of particular interest are the selector valve-switching valve combination, a core system for regulating a sequence of reactions, a core system adapted for polymer sequence analysis, and processes that implement the systems. The core systems can, as indicated by the use of the word "comprising" in their description, be either used as stand-alone systems or be part of larger systems.

Selector valve-switching valve combination

An important aspect of the invention is a rotary valve combination comprising a rotary selector valve (with its plurality of peripheral ports) connected by its central common port to a peripheral port of a rotary switching valve. The connection is preferably accomplished by a conduit means (e.g., a tube or capillary). It is preferred that the conduit means have an internal volume between 0.5 μ L and 10 μ L.

The rotary selector valves

It is preferred that the rotary selector valve comprise, in a first unit (e.g., the stator), a

central common port and a circular array of a plurality of peripheral ports such that the main axis of the common port is the same as the main axis through the circular array, and wherein said valve comprises, on a second unit, a radial connector channel, the first and second units being juxtaposed such that there is a single continuous selector channel formed by the common port, the connector channel, and a peripheral port, the selection of the peripheral port under control of the computer (the control preferentially effectuated by rotating the second unit), the internal volume of the continuous selector channel being the internal volume of the valve,

Preferably the two units are positioned in flat contact with each other, the flat contacting surface of the first unit against the flat contacting surface of the second unit. Contact between the two units is such that both the common port and a peripheral port meet the connector channel. This is optimally done by constructing the connector channel as a groove on the contacting surface of the second unit. It is preferred, but not required, that the common port act as an output port and that the connected peripheral port act as an input port, but the reverse relationship between the ports is also possible. The total internal volume (central port plus connector channel plus peripheral port) is preferably 1.5 μL or less (more preferably in the range 0.04 μL to 1.5 μL , most preferably 0.1 μL to 0.5 μL).

The rotary switching valves

It is preferred that the rotary switching valve comprise a circular array of three or more peripheral ports in a first unit (e.g., the stator) and a connector channel in a second unit, the two units being juxtaposed such that there is a continuous switching channel formed by a first peripheral port, the connector channel, and a second peripheral port, the selection of two peripheral ports being under the control of the computer (the control preferentially effectuated by rotating the second unit), the internal volume of the continuous switching channel being the internal volume of the valve.

Preferably the two units are positioned in flat contact with each other, the flat contacting surface of the first unit against the flat contacting surface of the second unit. Contact between the two units is such that two peripheral ports meet the connector channel. This is optimally done by constructing the connector channel as a groove on the contacting surface of the second unit. (The groove can be straight or curved). The total internal volume (central port plus connector channel plus peripheral port) is preferably 1.5 μL or less (more preferably, in the range 0.04 μL to 1.5 μL , most preferably 0.1 μL to 0.5 μL).

System for regulating a sequence of reactions

In a general aspect the invention is a computerized system for regulating complex reaction sequences in small volumes of solution, said system comprising:

a reaction vessel (for holding a volume of fluid in which a reaction takes place);

a first and second reagent vessel (for holding fluid reagents), said vessels each connected to a gas source,

a rotary valve combination connected to receive fluid from said reagent vessels and to deliver fluid to said reaction vessel, said combination optionally with other receiving-delivery capabilities, said rotary valve combination comprising a multi-position rotary selector valve connected to a rotary switching valve;

a computer connected to said valve combination so that said computer controls both whether the switching valve is configured to allow or to not allow fluid flow to the reaction vessel and whether the selector valve is configured for input from the first or second reagent vessel; and

wherein the internal volume of each of said valves along the path of fluid flow, is 1.5

μL or less, (preferably in the range $0.04 \mu\text{L}$ to $1.5 \mu\text{L}$, more preferably $0.1 \mu\text{L}$ to $0.5 \mu\text{L}$).

Each gas source is selected from a group of one or more gas sources each comprising a gas under pressure, said gas optionally differing from source to source. Any gas or gas combination that is not chemically reactive with a fluid or other reagent in the system can be used.

System adapted for polymer sequence analysis

In another aspect, the selector valve of the above-noted system is a first selector valve and the switching valve thereof is a first switching valve, and the system further comprises:

a second rotary selector valve connected to a second rotary switching valve;

a third rotary switching valve;

a conversion vessel connected to receive output from said third rotary switching valve,

a restraining means for restraining unprocessed polymer in the reaction vessel;

third and fourth reagent vessels connected to provide fluid to said conversion vessel;

wherein said second rotary valve, second switching valve, and third switching valve are each under the control of the computer and each have an internal volume of $1.5 \mu\text{L}$ or less (preferably $0.04 \mu\text{L}$ to $1.5 \mu\text{L}$, more preferably $0.1 \mu\text{L}$ to $0.5 \mu\text{L}$).

In particular embodiments, the system further comprises one or more of the following:

one or more wash solution vessels (each for holding a volume of fluid, preferably a wash solution), each said vessel connected to a gas source, each said vessel providing optional input (directly or via other elements of the system) via a selector valve to the reaction vessel and/or conversion vessel; and

a switching valve, under the control of the computer, for controlling fluid flow to an analytical device (such as an HPLC column or electrospray ionization mass spectrometer) from the conversion flask.

Reaction vessel and conversion vessel

It is preferred that, when the first and second fluid reagents are in the reaction vessel, the volume of fluid in the reaction vessel is in the range of 0.5 μL to 10 μL (preferably 1 μL to 5 μL).

It is preferred that a volume of fluid in the conversion vessel be in the range 1 μL to 150 μL (preferably 3 μL to 50 μL , most preferably 3 μL to 15 μL).

Selector valve cascade

An aspect of the invention is a rotary selector valve cascade in which a plurality of peripheral ports of a first selector valve are each connected to the central common port of one of plurality of rotary selector valves that each comprise a plurality of peripheral ports, the total internal volume of each valve being 1.5 μL or less (more preferably in the range of 0.4 μL to 1.5 μL , most preferably 0.1 μL to 0.5 μL).

Process aspect of the invention

In another general aspect, the invention is a reaction process under the control of a computer said process comprising the steps of:

- 1) Delivering, from a reagent vessel under pressure from a gas source, a volume of a reagent via a delivery line to a reaction vessel wherein the total volume delivered for purposes of a reaction in the reaction vessel is in the range 0.5 μL

to 10 uL, preferably 1 uL to 5 uL, and wherein flow through the delivery line is under the control of a computer-controlled rotary valve combination;

Preferred is the process with an additional steps (2) and (3):

- 2) Washing said valve combination and delivery line with a wash-solvent; and
- 3) Repeating step (1) optionally with a reagent, reagent vessel and/or gas source different from that used in step (1).

Process aspect of the invention used to sequence polymers

In an aspect of the invention adapted for sequencing a polymer, the process comprises the steps of:

- (1) Placing a polymer (preferably in the range 5 to 500 femtomoles, more preferably 20 to 250 femtomoles, most preferably 50 to 200 femtomoles) in a reaction vessel, said polymer restrained in said vessel, the restraining preferably accomplished by adsorption of the polymer to a solid support;
- (2) Delivering to said reaction vessel, from a first reagent vessel under pressure from a gas source, a volume of a first fluid, said fluid comprising a first reagent, which reagent will react with a terminal monomer on the polymer;
- (3) Delivering to said reaction vessel, from a second reagent vessel under pressure from a gas source, a volume of a second fluid, which fluid (the fluid can be a solvent or a solution of a solute in a solvent) will cause a terminal monomer-reagent moiety to be cleaved from the polymer;
- (4) causing all or part of the fluid in the reaction flask to be transferred under gas pressure via a delivery line to a conversion flask while allowing the polymer, less the terminal monomer, to remain in the reaction vessel;

- (5) Prior or after step (4), delivering from a third reagent vessel under pressure from a gas source a volume of a third fluid such that the combination of said fluid and the fluid transferred in step (4) to the reaction flask will cause the terminal monomer-reagent moiety to be cleaved to create a terminal monomer free of covalently bound reagent;
- (6) Transferring under gas pressure from a gas source said terminal monomer created in step (5) to an analytical device that will identify the nature of the terminal monomer.

It is preferred that each volume delivered in steps (2), (3), (4), (5), is in the range 0.2 uL to 10 uL, (preferably 0.5 uL to 5 uL). Independently, it is preferred that the volume delivered in step (6) is in the range 1 uL to 10 uL, (preferably 2 uL to 5 uL).

In a preferred embodiment of the process adapted for sequencing a polymer, the process comprising steps (1) - (6), is performed a plurality of times.

Air-free reaction system

In preferred embodiments of the processes, the reaction vessel and the conversion vessel are kept free of air and oxygen. The ability to do this effectively is an advantage of the present invention.

Sequential treatment with acid and base

In particular embodiments of the processes, the process comprises sequentially delivering acid, organic solvent, and base to the reaction vessel. The ability to do this effectively is an advantage of the present invention.

Applications of the invention

Examples of such applications are:

- 1) automated reactor for solution chemistry;
- 2) automated reactor for liquid "flow-through" chemistry;
- 3) automated reactor for gas-phase chemistry;
- 4) automated sample preparation, modification, and reaction prior to electrospray ionization mass spectrometry;
- 5) automated enzymatic digestion of proteins, nucleic acids and complex carbohydrates;
- 6) thermostatic control for sample preparation, modification and reaction (including the above uses);
- 7) automatic precision volume metering utilizing loops with volume range 0.1 microliter and above;
- 8) automated selection of samples for any use described above.

Specific polymer applications

In one set of embodiments, the system is adopted for processing a polymer in step with fashion, one monomer at a time.

Example of such processing are:

- 1) the determination of the amino acid sequence of a polypeptide;
- 2) the determination of the base sequence of a nucleic acid; and
- 3) the determination of the sugar sequence of a polysaccharide;

Brief Description of the Drawings

Figure 1. (A) Schematic diagram of the 'PSMSK' automated microfluidics system. The instrument **100** is shown here in a protein sequencer configuration. Connections of the computer **17** to the other components of the system are not shown but are described in the text. (B) Legend to selected symbols seen in (A). (C) Examples of materials used with system shown in (A).

Figure 2. PSMSK 'function #29', *deliver liquid TFA to cartridge*. This function consists of the seven sequential 'events', i.e. unique valve position combinations, as shown in panels A-G and listed in panel H where events 1-7 correspond to panels A-G, respectively. Only rotary valves **51**, **61**, **66** are shown; for the precise positioning in the entire system and for the key to symbols and abbreviations, see schematic in Fig. 1. Valve **51** is a ten-port selection and valves **61**, **66** are two-way switching (A/B); solenoid valves (not shown) are used to pressurize or vent the TFA bottle (ON/OFF). The flow path through the valves during each event is shown in bold; no liquid or gas flow occurs during 'event #7'. Total function time is 1 min 44 sec. 'Function #29' is used in the reaction cycle (Table 2A) as **step 30**.

Figure 3. 'On-line' capillary-LC analysis of 25 fmol (delivered to the flask) of PTH-amino acid standard under conditions as described in the text. A 300-micron i.d. column, packed with C₁₈ PTH (5-micron particle size) resin, was used at a flow of 3 μ L/min; absorbance detection was done at 269 nm. Only the relevant portion of the chromatogram is shown. More details can be found in the 'Materials and Methods' section.

Figure 4. Micro-chemical sequencing of 100 fmol bovine β -lactoglobulin using the 'PSMSK' microfluidics system and on-line, capillary-LC PTH-amino acid analysis.

Experimental conditions were as listed under 'Materials and Methods' and in the **Figure 3** legend. Chromatograms 1-16 are shown; absorbance detection full scale varies, as indicated. PTH-amino acid peaks are indicated with an arrow and the femtomole amounts (corrected for background) shown.

Figure 5. Schematic drawing of selected portions of a selector valve. A. A first side planar view. B. A sectional view. C. A second side planar view.

Figure 6. Schematic drawing of selected portions of a switching valve. A. A first side planar view. B. A sectional view. C. A second side planar view.

Figure 7. Top Block of reaction vessel. A. A planar view. B. A sectional drawing along the line B'-B' in A. C. A bottom planar view.

Figure 8. Bottom block of reaction vessel. A. A bottom planar view. B. A sectional drawing along the line B'-B' in A.. C. A top planar view.

Figure 9. Schematic sectional view of reaction vessel chamber and contents created by combining the bottom block (**Fig 8**) and top block (**Fig 7**) of the reaction vessel with additional components.

Figure 10. Schematic partially sectional drawing of selected portions of a conversion flask.

Figure 11. Schematic partially sectional view drawing of a selected portion of a rotary valve combination in which a valve (**Fig. 6**) is connected via a capillary to a selector valve (**Fig. 5**).

Detailed Description of the Invention

Abbreviations and glossary

ATZ-aa stands for anilino thiazolinone amino acid.

BLG stands for beta-lactoglobulin.

BuCl stands for butyl chloride.

CLC stands for capillary liquid chromatography

A computer is an electronic device that receives input information in electronic form and processes that information electronically or electromagnetically to create electronic output signals that can be converted to printed material, converted electronically to displayed material, or otherwise further processed. Typical computers are personal computers such as made by IBM or other companies but both more sophisticated computers can also be used and, conversely, the computer can be one of much simpler construction than a personal computer by virtue of the fact that it is dedicated to the process and systems of the invention.

CZE stands for capillary zone electrophoresis.

DMPTU stands for dimethylphenylthiourea.

DPTU stands for diphenylthiourea

EtAc stands for ethyl acetate.

GP stands for gas-phase.

HPLC stands for high performance liquid chromatography.

LC stands for liquid chromatography.

Lys stands for lysine.

10% MeCN and 10% MeCN/H₂O stands for 10% acetonitrile in water.

MS/MS stands for tandem mass spectrometry.

ESI stands for electrospray ionization.

PCR stands for polymeric chain reaction.

PITC stands for phenyl isothiocyanate.

Plurality means 2 or more.

PSMSK stands for the automated micro-fluidics system 100.

PTC stands for phenyl thiocarbamyl.

PTH-aa stands for phenyl thiohydantoin amino acid.

Reagents include any fluids or solids that participate in chemical reactions and also any solution or solution components in which said fluids or solids are dissolved or suspended.

RP-HPLC stands for reversed-phase high performance liquid chromatography

25% TFA and 25% TFA/H₂O stands for 25% TFA (v/v) in H₂O

0.1% TFA stands for 0.1% TFA in H₂O

TFA stands for trifluoroacetic acid.

THF stands for tetrahydrofuran.

TMA stands for trimethylamine.

The terms "path of fluid flow" and "flow through channel" are used interchangeably.

For a particular component of the system in a given configuration (e.g., a particular valve setting), the terms represent the total path followed by fluid within that component in that configuration.

"Under pressure" means under greater than atmospheric pressure.

Val stands for valine.

A "volumetric loop" is a conduit that can be either straight, looped, helical, or otherwise curved. "Loop" here is a term of art.

"Wash solutions", also referred to as "wash solvents", are solutions that are used to

wash a vessel or a vessel-to-vessel connector after it has been vacated by a reagent.

Overview of the invention in relation to the Figures

The following, by relating the invention to the Figures, is intended to exemplify the invention, not limit it.

Figure 1A shows a computerized system **100** for regulating complex reaction sequences in small volumes of solution. The system contains a reaction vessel **71** for holding a volume of fluid **81** and reagent vessels **1-6** (for holding fluid reagents **11-16**, respectively). The vessels **1-6** are each connected to a gas source, selected from a group **171**. Gas sources **171a-f** each contain a gas **174a-f** under pressure. The system also has a rotary valve combination **59** connected to receive fluid from reagent vessels **1-2** and to deliver fluid to the reaction vessel **71**. The combination can be seen to have other receiving-delivery capabilities. The rotary valve combination **59** comprises a multi-position rotary selector valve **51** connected via a conduit means **121** to a rotary switching valve **61**. A computer **17** is connected to the valve combination **59** (Computer connections are described in the text herein but are not shown in the Figures.) The computer controls both whether the switching valve **61** is configured to allow or to not allow fluid flow to the reaction vessel **71** and whether the selector valve **51** is configured for input from reagent vessel **1** or from reagent vessel **2**.

The system is adapted for polymer sequence analysis. To that purpose it includes a conversion vessel **101** for holding a volume of fluid **111**. It also includes a restraining means **191** for restraining the unprocessed portion of the polymer in the reaction vessel **71** (See **Figure 7**). The restraining means can, as in the example below, provide a surface to which

the polymer can noncovalently adsorb, or it can be a component to which the polymer, e.g., via an end group, can be covalently linked. The system also has a rotary switching valve 62, under the control of computer 17, for controlling fluid flow to the conversion vessel 101 from the reaction vessel 71. The system also has a second rotary valve combination 58 connected to receive fluid from reagent vessels 4-5 and to deliver fluid to flow to said valve 62. The combination can be seen to have other receiving-delivery capabilities. The rotary valve combination 58 comprises a multi-position rotary selector valve 52 connected to a rotary switching valve 65. The computer 17 is connected via a conduit means 121 to valve combination 58 so that said computer controls both whether the switching valve 65 is configured to allow or to not allow fluid flow and whether the selector valve 52 is configured for input from reagent vessel 4 or from reagent vessel 5.

The system has additional components. It has wash solution vessels 41-44 (for holding wash solutions 141-144), the vessels connected to a gas source selected from group 171. Vessels 41-43 are connected as optional input to valve combination 59. Vessel 44 is optional input to valve combination 58. Additionally, the system has an analytical device 151, an HPLC column in combination with a UV monitoring device. Other features of interest are a switching valve 64 under the control of computer 17, for controlling fluid flow to the HPLC column from the conversion flask 101. Reagent vessels 221 and 222 are sources of reagents 181 and 182 for the HPLC column. Waste vessels 201 collect waste fluid.

Multiposition rotary selector valve 51 is schematically shown in Figure 5 and, in the first unit 249 of the valve, comprises a central common port 241 and a circular array of a plurality of peripheral ports 231a-j, (only 231a and 231f are marked in Figure 5) such that the main axis (imaginary line 135) of the central common port is the same as the main axis

through the circular array. The valve also comprises, on a second valve unit 245, a radial connector channel 251 in a second unit 245 of the valve. A continuous selector channel is created by the common port 241 plus the conduit means 251 plus the peripheral port 231a. The common port 241 can be connected to any peripheral 231a-j port selected by rotation of the second unit as determined by the computer 17.

Rotary switching valve 61 is schematically shown in Figure 6. The valve comprises, in a first valve unit 283, a circular array of four peripheral ports 271a-271d and, in a second valve unit 246, further comprises connector channels 281 and 282. In Figure 6, a continuous switching channel is created by peripheral port 271b plus connector channel 281 plus peripheral port 271a. A continuous switching channel is also created by peripheral port 271c, plus connector channel 282, plus peripheral port 271d. The second unit 246 is rotatable about its main axis (imaginary line 136) under the control of the computer 17, so that connector channel 281 can form a continuous switching channel with, peripheral ports 271a and 271d. In the latter situation, channel 282 forms a continuous switching channel with peripheral ports 271b and 271c.

A rotary valve combination 59 comprising a rotary selector valve 51 connected by its common central port 241 via a conduit means 121 (e.g., a tube or capillary with an internal passage 122) to the peripheral port 271c of a rotary switching valve 61 is shown schematically in Fig. 11..

EXAMPLE

Materials

Here, we report development of a novel instrument with miniaturized reaction

cartridge and conversion flask (each about 1/10th the size of a high-end commercial instrument -- Applied Biosystem's model 494cLC) and reagent / solvent flow paths. Our sequencer allows identification of amino acid derivatives well below 50 femtomoles, with repetitive yields in the 90-94% range; chemical background is very low. Because the instrument incorporates a miniature, pressurized flask with glass capillary pick-up line, it could conceivably be interfaced with continuous-flow, nanospray-type mass spectrometry, as recently reported,³¹ for mass analysis of any peptide or amino acid derivatives. The position of the mass spectrometer would be the same as the HPLC column in **Figure 1**.

All reagents and solvents (see also **Fig. 1**), PTH-amino acid standard, β -lactoglobulin standard, polybrene ('biobrene'), glass fibre discs, and cartridge seals were obtained from Applied Biosystems (Foster City, CA), except where noted. Acetonitrile was from Burdick & Jackson (Muskegon, MI), PTH-norleucine from Pierce (Rockford, IL), and the highest purity argon and helium gas from TechAir (White Plains, NY).

Microfluidics Sytem: Edman Chemistry Configuration

The automated micro-fluidics system **100** (also referred to as the "PSMSK sequencer" or simply "PSMSK") can be better understood in relation to **Figure 1**. The system is built around an array of eight rotary valves, connected with fused silica capillary (50, 75, 100, and 150- μ m I.D. /365- μ m O.D.; Polymicro Technologies; Phoenix, AZ) to two miniaturized reactors, a reaction cartridge (functioning as a reaction vessel) **71** and a conversion flask (or conversion vessel) **101** (see **Fig. 1**). The system has an extremely low internal volume of all components in the fluid path (see **Table 1**), which makes possible the automated precision metering and delivery of reagents in the sub-microliter range.

Cheminert series (Valco, Houston, TX) rotary valves were chosen for the fluid system because of their low internal volumes, small size, and ease of computerized control. These valves have 150- μ m port diameters and a total internal volume of less than 0.4 μ L. By mounting the

valve rotor seal with a lower torque, greater longevity is obtained at a sealing pressure of 250 psi (instead of a normal factory-rated 5,000 psi). The wetted surfaces are fabricated out of the inert polymers, PAEK and Valcon E. PEEK nuts and one-piece fused silica adapter ferrules (Valco #ZN1FPK-10, ZF1PK) are used to connect fused silica capillary to the valve ports. Two basic categories of rotary valves are used in the system. The multi-position selector valve is characterized by a central common port and a circular array of peripheral ports positioned at the outer circumference of the valve stator. A rotor seal 245 with a single radial fluid path connects the central common port to one of the peripheral ports, determined by the angle of rotation. Ten-port valves 52 and 51 of this type (Figure 5) are used as a conversion selector valve 52 and a reaction selector valve 51 respectively to select reagents and argon gas for delivery to the conversion flask and reaction cartridge, respectively (Fig. 1). Two-position switching rotary valves have ports positioned around the outer circumference of the stator only (Fig. 6). The rotor seal 246 in a switching valve typically has more than one notch and connects adjacent ports in an arc shaped path. Six-port switching valves 64 and 63 are used as an injector valve 64 and a flask valve 63. Four-port switching valves 61, 62, 65, 66 are used as transfer valve 62, conversion switching valve 65, TFA isolation valve 66, and reaction switching valve 61 (See Fig. 6). To create the specialized flow paths for Edman chemical sequencing and to eliminate any dead volumes in the system, many of the switching valves incorporate rotors (rotor seals) that are custom fabricated as to their number of fluid paths. (Valco, Houston, TX). Valves 65, 66 and 61 have rotors with only one fluid path instead of the standard two; the rotor of valve 63 has two fluid paths instead of the standard three. Ports that are unused are plugged with PEEK nuts and solid Teflon tubing (Zeus, Orangeburg, SC).

By connecting the common port of a selector valve via a capillary to a peripheral port of a switching valve, the connecting capillary may serve as a precision metering loop. (Figure 11) This arrangement is used to deliver consistent volumes of reagents to both the reaction cartridge and the conversion flask 101. For instance, the reaction selector valve 51 is coupled to the

reaction valve **61** with a section of capillary that forms a 0.5- μ L loop **121**, used to meter liquid TFA and PITC to the reaction cartridge **71** (**Figs. 1 and 11**) . The conversion selector valve **52** is coupled to the conversion switching valve **65** with a section of teflon tubing forming a 7.5- μ L loop **124**, used to deliver a metered volume of PTH-amino acid standard, 25%TFA, and 10%MeCN to the flask **101** (**Fig. 1**). It is especially useful for the consistent injection of sample to the HPLC system. The use of this metering loop is described further below.

Each valve has its own integral microelectric actuator for precise control of rotation with a 100-msec switching time (actuator not shown in the Figures). The instrument control software sends simple ASCII commands via standard RS-232 serial communication to control the actuator position. Each valve is programmed with its own unique identity number. The commands are prefaced with this I.D. tag, allowing the control of several valves simultaneously over one daisy-chained serial connection. Since the valve rotation is bi-directional, the valves are initialized by the software to take the shortest possible rotation path when switching between ports. This feature is considered strategically when programming valve functions to ensure that certain chemicals (i.e. TFA and TMA) never mix (see **Figure 2**).

The current system includes twelve 40-mL bottles **1-6, 41-44, 181-2** for a wide range of customized chemistries. Each bottle is pressurized with argon gas through an inert teflon solenoid valve (Angar Scientific; Cedar Knolls, NJ) controlled by logic signals from the instrument control software of a computer **17**. Inert PEEK check valves (Lee Company, Westbrook, CT) are fitted on the bottle pressure lines to ensure that chemicals do not contaminate the argon regulation system.

There are six argon regulators **171a-171f** that are in gas sources and which provide argon at pressures of 2.0, 2.0, 4.0, 12.0, 24.0 and 50.0 psi, respectively. (Porter Company, Hatfield, PA) (see **Figure 1**). Two are dedicated to pressurizing the TFA bottle **3** and TMA bottle **2**; the remaining four are available for supplying the other bottle positions, for providing varying levels of pressurized argon to deliver reagents to the reactors, and for drying operations. The pressure

of each regulator is monitored by a chip-based pressure transducer (SenSym, Milpitas, CA) and displayed in the instrument status window on the computer.

Reaction Cartridge

There are two reactors in the system when configured for Edman chemistry, a flow-through 'reaction cartridge' 71 and a solution chemistry 'conversion flask' 101 (Fig. 1). Various features of the reaction cartridge are shown schematically in Figs. 7, 8 and 9. The reaction cartridge 71 consists of two 0.5-inch diameter /0.5-inch high cylindrical blocks 233 (Fig. 7) and 234 (Fig. 8), manufactured out of borosilicate glass (United Lens; Southbridge, MA), stacked vertically to form the reaction chamber. Both blocks were bored with a 0.015-inch fluid path 225 through the central axis. One flat face 226, 227 of each block was machined with a 0.25-inch cylindrical cavity 237, 238 for a pressure fit tubing connection utilizing a ferrule. The second flat face 224 of the bottom block 234 was machined with a conical cavity 226 contiguous with a 3-mm circular cavity 236 to hold a glass fiber support membrane 191 for immobilizing the protein (Figure 7). The second face 228 of the top block 233 was machined with a complimentary mating surface including a rim 239 that seals precisely around the circumference of the glass fiber disc with the use of a permeable Teflon seal 235 (See Fig. 9). The second face 228 was also machined to include a conical cavity 242.

The cartridge 71 is assembled in an aluminum heater (not shown) that is thermostatically regulated by the instrument control software. The heater has a window and neon back lighting to view the contents of the cartridge. The top block 233 is held captive inside the reaction cartridge assembly at all times. It is only removed for occasional cleaning. For sample loading, the bottom cartridge block 234 is removed from the assembly in a stainless steel holder with a simple "half-twist" mechanism. Once the protein 195 has been pipetted on the disc 191, a permeable teflon seal 235 is placed on top of the bottom block 234 (See Fig. 9). The cartridge 71 is then fully assembled by mounting the stainless steel holder (not shown) to the heater with

a second half-twist. A spring steel bellville washer is located at the top of the cartridge to ensure a tight seal while protecting the glass blocks from damage during the assembly. The cartridge is pressure tested prior to all sequencing runs to ensure that the reaction chamber is an isolated, oxygen-free environment, minimizing background artifacts produced during the coupling and cleavage reactions.

There are two rotary valves **61** and **62** that control flow through the reaction cartridge. Located at the bottom of the cartridge is the reaction switching valve **61**, that controls delivery of reagents and argon gas from the reaction stream selector valve. The direction of flow during the sequencing chemistry is from the bottom upward through the sample carrier. The transfer valve **62** at the top of the reaction cartridge is a two position switching valve, that directs the outlet of the cartridge to the conversion flask for further processing or to the waste bottle.

Conversion Flask

The conversion flask **101**, illustrated schematically in **Fig. 10**, is a 75- μ L borosilicate tapered glass vial (Pierce; Rockport, IL). It is mounted inside an aluminum heater (not shown) thermostatically regulated under the control of the computer **17**. Tubing connections are made to the flask through a teflon seal **102** at the top. The vial can be removed easily from the heater for cleaning or exchange by unthreading a stainless steel thumbscrew located at the bottom of the assembly. There are two tubing connections **131**, **132** to the flask **101**. The first is a vent line **131** made from 1/16-inch O.D./0.004-inch I.D. teflon tubing. It is installed with a simple pull-through friction fit through a 0.056-inch hole in the teflon seal. The vent line is cut flush with the inside of the teflon seal using a razor blade. The second tubing connection to the flask is a pickup line **132** made out of 75- μ m I.D. / 365- μ m O.D. polyimide coated fused silica capillary. To install the pickup line first a 0.01-inch I.D. / 0.0625-inch O.D. teflon tubing sleeve **137** is pulled through a 0.056-inch hole in the teflon seal at the top. This sleeve is cut off flush with the inside of the teflon seal using a razor blade. By pulling the capillary **132** through the teflon sleeve a pressure

tight seal is formed. A long enough section of capillary is pulled through the teflon seal so that it reaches the bottom of the flask. A bunsen burner is used to burn off the polyimide coating from the portion of the pickup line that is exposed to reagents, as it was found that the polyimide breaks down during the conversion process and interferes with the liquid chromatography.

A 6-position, 2-port switching valve **63** (see **Figure 1**) is located above the flask **101** and controls the delivery of fluids and argon gas. In position A, the vent is opened to waste and the pickup line **132** is connected to the transfer valve **62**. This position is used to deliver fluids to the flask **101** in a controlled fashion from the bottom up, and also to bubble argon through the sample. This argon dry operation is used in many situations, including the conversion of ATZ-amino acids to their PTH counterparts, solvent evaporation prior to HPLC injection, and high pressure flask cleaning between cycles. In position B, the vent line **131** is connected to the transfer valve **62** and the pickup line **132** is connected to the injector valve **64**. This position is used to empty the contents of the flask **101** to the injector loop **123** or the waste bottle **201**. By delivering pressurized argon from the reaction selector valve **51** to the vent line **131** the contents of the flask **101** are forced out through the pickup line **132**.

A 10-cm long, 50- μ m I.D. restrictor capillary **134** is connected to the outlet of the injector valve **64**. The narrow internal diameter of the restrictor capillary provides sufficient back pressure to maintain a gradual 15 second loading of the injector loop **123** when argon at 12.0 psi from gas source **171d** is used to pressurize the flask. The low internal volume of the restrictor capillary is also conducive to a high injection percentage. When the first sign of liquid is present at the end of the restrictor capillary, only 0.2- μ L of sample has passed beyond the loop. This makes it possible to routinely inject 5 μ L out of a sample volume of 7.5 μ L.

HPLC System

PTH-amino acids are separated on a 25-cm long, 300- μ m I.D. capillary

HPLC column, **151** custom packed (LC Packings, San Francisco, CA) with 5- μ m C₁₈ PTH stationary phase. The column **151** is placed inside a heated enclosure (Eppendorf; Westbury, NY)) and thermostatically regulated at 55°C. HPLC solvent A **221** is 5% tetrahydrofuran in water, with 20 mL of 'Premix Buffer' added per liter (Applied Biosystems); solvent B **222** is acetonitrile (Burdick & Jackson) with 500 nmol of dimethylphenylthiourea (DMPTU; Applied Biosystems) added per liter. Solvents are contained in Kontes Ultraware (Vineland, NJ) reservoirs **181** and **182** and are sparged with helium gas at a pressure of 3 psi. The gradient is formed by an Applied Biosystems 140D micropump at a flow rate of 30 μ L/min; a flow splitter (LC Packings) with a 10:1 split ratio delivers a consistent gradient (12-24%B/ 4 min; 24-50%B/ 22min; 50-90%B/ 1 min) to the column at a flow rate of 3 μ L / min. The flow splitter **232** also contains a micro-volume union and static mixer. The outlet of the column is directly connected to a UZ-View 'ball lens' capillary flow cell (LC Packings) which is mounted in an Applied Biosystems model 785A UV absorbance detector. All connections in the LC system are made with 75- μ m I.D. / 365- μ m O.D. fused silica capillary. The detector is set to a wavelength of 269 nm, with a rise time filter setting of 2 sec. The absorbance data is collected from the UV detector by a Nelson 900 Series analog to digital converter, filtered, stored, and analyzed using TurboChrom 4 software (PE Nelson; Cupertino, CA).

Instrument Control

The instrument **100** is controlled by a computer **17**, a Macintosh 225MHz computer (Apple; Cupertino, CA) comprising and running a software application, which was programmed in the LabView environment, version 5.1 (National Instruments, Houston, TX). A DaqCard 1200 (National Instruments) data acquisition and control card is installed in the computer and is used to interface between the software and circuit board contained in the instrument enclosure. The DAQ card has 8 analog to digital (A-D) converters and 24 lines of digital input/output. The A-D converters monitor the signals from the reaction cartridge and conversion flask temperature

sensors as well as the argon regulator pressure transducers. The digital lines control the solenoid valves, reaction cartridge and conversion flask heater elements, and four relays used to control external devices (e.g. HPLC pump, UV detector, and data collection system). The rotary valves are controlled by serial communication directly from the computer's serial port. Communication from the computer to the sequencer is accomplished through one DB-9 RS-232 serial cable and one 50 conductor ribbon cable. These connect to ports on the back panel of the instrument. Also located on the back panel are a cooling fan, bulkhead connection for the waste bottle vent line, and the four relay outputs for controlling external devices.

The circuit board is multifunctional. It buffers the low millivolt signals from the pressure transducers and temperature sensors and converts them to the useable range of the DAQ card A-D converter. The 5VDC digital logic signals from the DAQ card trigger transistors on the circuit board to switch higher current power to the solenoid valves, relays, and heater elements. In the case of the solenoid drivers, an additional circuit drops the initial 24VDC 'strike' voltage down to 12VDC after 100 msec. The board also acts as a voltage regulator, supplying the correct power to the instrument's cooling fan and neon lighting. The circuit board was created from a copper clad board with a 'photo-resist'" coating (Kepro Circuit Systems, St. Louis, MO). A negative of conductor circuitry, designed with CAD software, was printed on a transparency, and placed on the board, after which the board was exposed to UV light in a darkroom. Hot sulfuric acid was used to etch away unneeded copper, leaving the desired conductor circuitry.

The Lab View software environment allowed the design of a graphical display that emulated the actual operation of the instrument. The status of all valves, relays, regulator pressures, and temperature transducers are displayed on the monitor screen 18 in real time as the instrument is operated. The software has two modes of operation, manual and automatic. In both modes the same real time graphical display is utilized. In the manual mode a manual control window is also present on the computer desktop. This window contains buttons for the control of every piece of hardware at the user's discretion. In the automatic mode, control cycles are

preprogrammed by the user. In this case, the manual control window is replaced by a sequencer status window that displays information regarding the current sequence, cycle, and function being performed.

An automated control program is created through the use of user-defined reaction and conversion functions (see **Table 2**). A sequencer cycle is one complete program for analyzing one amino acid residue of the protein. Distinct reaction and conversion cycles are created independently to control the activity of the two distinct reactors **71**, **101**. The software then meshes these two cycles together automatically so that the transfer of sample between the reaction cartridge **71** and conversion flask **101** is coordinated properly. Several cycles are looped together to create a sequence program that can process as many amino acid residues as the user requires. Sequencer functions are not necessarily single command events. A function can include as many independent timed 'events' as needed. In fact, nearly all of the functions we used were more than one event, with a maximum of eight events. This feature is especially useful since most of the chemistry is automated with the use of metering loops. Delivering a loop of reagent is a multi-step procedure, as illustrated by the metered delivery of TFA liquid **13**, the most complex case, in **Figure 2**.

Operation

The reaction cartridge **71** was designed with a 3-mm diameter disc **191** to reduce chemical and amino acid background, a requirement to call amino acid sequence from the chromatographic data. Yet, the size should be large enough to allow for a practical sample loading volume. At a diameter of 3 mm, a glass fiber disc can hold a sample volume of 2 μ L. Several 2- μ L loads to a single filter can be done in series to accommodate samples of larger volume. Prior to sample loading, the glass fiber disc **191** is coated with 0.18 mg of polybrene, a polymeric sample carrier that increases the protein affinity for the glass fiber. The unloaded, polybrened disc is then precycled by running several standard sequencing cycles until the

chromatogram appears clean. BLG protein standard was pipetted onto the disc, dried using 2-psi argon gas, and the reaction cartridge was reassembled and pressure tested. The reaction and conversion cycles (**Table 2**) were initially based on the custom cycles used with commercially available sequencers in our lab.¹⁷ They have since been extensively modified to meet the requirements of the new miniaturized flow path and adjusted according to experimental observations.

Table 2A illustrates in detail reaction cycle #27 "Glass Fiber Disc". **Steps 1-11** comprise the coupling of PITC **11** to the N-terminal amino acid of the protein. TMA vapor **12a** originating at 2 psi from solution **12** is delivered three distinct times, at 400 seconds each, for a total of 1200 seconds. TMA provides the basic environment necessary for proper coupling. Deliveries of 1- μ L aliquots of 1% PITC **11** are interspersed between the TMA steps. The overall coupling time in the reaction cycle was set intentionally long to ensure minimal sequencing lag that would otherwise result in inefficient coupling chemistry. This is then followed by a series of solvent washes (**Steps 12-29**) with heptane **141**, ethyl acetate **142**, and butyl chloride **143**. The volume of the washes was optimized to minimize chemical background while preventing sample washout (see **Table 1**). For subsequent acid cleavage (**Step 30**), delivery of TFA **13** was also tightly controlled, as to achieve low lag while minimizing sample washout. We found that a volume of 0.5 μ L was most effective. This step is described in detail in **Figure 2**. Transfer of ATZ-amino acids to the conversion flask **101** (**Steps 36-55**) was the subject of rigorous optimization. Since the volume of the flask is only 75 μ L, solvent volume should balance efficient extraction of the ATZ-amino acids with this limitation. Thus, the delivery of the transfer solvent (BuCl **143**) was broken up into several smaller steps, with argon dries interspersed between, to keep the solvent in the flask at a manageable level. This procedure also increases the efficiency of the extraction, as shown.¹⁷ Following the transfer phase, the reaction valves are cleaned with BuCl, **143**, Heptane **141**, and EtAc **142**, to prepare them for the next amino acid cycle (**Steps 58-61**).

Table 2B details the conversion cycle utilized, #28 "Low Pressure Conversion". It begins with an extensive cleaning of the flask **101** (Steps **1-10**) with 25%TFA/H₂O **14** and 10%MeCN/H₂O **144** to eliminate any carryover between cycles. The cleaning solvents are delivered through the vent line **131**, instead of through the pickup line **132** as is the case with other deliveries during the sequencing run, ensuring better cleaning of all fluid pathways. Once filled with 40 μ L solvent, the highest available argon pressure (50 psi) from gas source **171f** is bubbled through the pickup line **111 + 132**/lower section to vigorously wash the entire internal volume of the flask, including the top teflon seal and the highest reaches of the vial. The flask valve **63** then switches position so that high pressure argon is delivered to the vent **131**, forcing the contents of the flask **101** to the injector waste **201**. To expedite the large volume deliveries of 25%TFA/water **14** and 10%MeCN/water **144** through the vent line, these bottles are pressurized with argon at 12.0 psi from source **171d**. After cleaning, a 225-femtomole, 7.5- μ L aliquot of PTH-norleucine **16** is added to the flask and evaporated (Steps **11-12**). Then, a 7.5- μ L loop load of the conversion reagent, 25%TFA/water **14**, is delivered to the flask **101** (Steps **13-14**), just before the transfer of ATZ-amino acid from the reaction cartridge **71** (Step **15**). Argon at 12.0 psi is bubbled through the flask **101** for better conversion reaction kinetics and to slowly evaporate off the 25%TFA (Step **16**). Once the flask **101** is completely dry, the injector loop **123** is flushed with argon to prepare for injection (Step **17**), and the contents of the flask **101** reconstituted with a 7.5- μ L loop of 10%MeCN, and bubbled with argon from source **171e** at 24.0 psi to improve dissolution (Steps **18-22**). Argon at 12.0 psi is then delivered to the vent line **131**, displacing the sample into the injector loop **123** (Step **23**). After 15 seconds, the injector valve **64** switches position, injecting the sample onto the column **151** and a relay closes to start the HPLC gradient program (Steps **24-25**).

RESULTS AND DISCUSSION

Proteins can be prepared, fractionated and detected at the femtomole level.

Taking into account a 2 to 5-fold loss during sample handling, chemical structure determination must then be done with just 100-200 femtomoles of starting material. Our objective, therefore, was development of an automated protein sequencer whose range of operation would include that range. We sought to do so by further miniaturizing the entire fluidics system, including reaction vessels and the analytical component, capillary (300-micron ID) LC, all in a very proximate arrangement.

Preliminary Considerations

We first constructed a stand alone, modular capillary-LC system (see **Materials and Methods**). Flow rates, gradients and column temperatures were optimized for various column/solvent combinations as to yield the best possible resolution, highest signal/noise ratio (in general and for each PTH-aa in specific) and least baseline drift. In anticipation of 'on-line' sample injections from a sequencer, we studied tolerance for increasing MeCN concentrations (3 to 10%) in function of progressively larger injection volumes (0.5 to 20 μ L). As 'sequencer-injected' samples may also contain trace amounts of TFA, its effects were also investigated. Initial attempts at optimizing operational parameters, injecting standards (20-100 femtomoles each of all PTH-aa) in an 0.5- μ L volume of 3% MeCN/0.1% TFA, resulted in complete separation. Most PTH-aa's can be detected at the 20 femtomole level, with signal/noise ratios better than 1 to 5, and injections of up to 5-8 μ L in volume appeared feasible (data not shown). Consequently, if analysis of 60-70% or more of the PTH-aa's after each cycle is desired, the analytes should be dissolved in less than 7-10 μ L of solvent, in turn necessitating a smaller conversion flask; e.g. 75 μ L, as compared to 750 μ L in current commercial sequencers. Space constraints required the use of 75- μ m I.D. / 365- μ m O.D.

capillaries for liquid delivery and pick-up. To avoid overfilling the smaller flask 71 during ATZ-aa transfer, the reaction disc 191 had to be reduced in size accordingly, to 3-mm diameter. Further reduction in wetted surfaces was accomplished by bringing chemistry vessels and detection system in close proximity, plumbed together with 50 to 100-micron ID capillary tubing as well, to create an integrated micro-fluidics system. Capillary bore tubing requires substantially higher pressures for delivering and drying liquids. Using a bread board instrument, it was found that increased back pressures cause liquids to back up in the wrong ends of the Z-path type valve blocks of an Applied Biosystems instrument. It was therefore decided to replace them with nanoliter-dead volume rotary valves in the final design, which also permitted accurate metering of reagents in the sub-microliter range.

Design and Construction

The instrument was constructed to conform to the stipulations summarized above and following the schematic diagram shown in **Figure 1**. It features the incorporation of 8 rotary valves (numbered in **Fig. 1**). The reaction cartridge 71 is positioned between valves 61 and 62 (liquid and gas flow from the bottom up) and the flask 101 right below valve 63. Note that, together, valves 51 (10-port selection) plus 61 (two-way switching) accomplish the same result as the "cartridge reagent/solvent-blocks" in an Applied Biosystems automated sequencer (see also **Fig. 2**); similarly, valves 65 and 52 accomplish the same result as the "flask reagent-block" in an Applied Biosystems automated sequencer. Valve 66 (**Figs. 1, 3**) serves to isolate TFA from the rest of the chemicals and solvents to exclude potential chemical problems, including salt formation upon contact with base (TMA). Rotary valves were selected on the basis of the lowest-dead volume and compatibility with TFA. Acid was continuously pumped through for an entire week without evidence of pressure leaks or visual damage. Cartridge 71, flask 101 and LC-injector valve 64 plus column 151 are all plumbed with glass capillary tubing; the rest with teflon capillary tubing.

The level of miniaturization and versatility incorporated in the design of our automated micro-fluidics system, as described here in the PSMSK sequencer, is best illustrated by comparison of the chemistry modules, selected wetted surfaces and variable pressurization to those in a state-of-the-art, commercially available sequencer (Applied Biosystems model "cLC494"). As shown in **Table 1**, the PSMSK reaction disc/chamber is about 4-8 times smaller, and eightfold less reagent and considerably less solvent are consumed; however, wash solvent per disc surface area is comparable. The overall flow path volume(E-block to injector) is 30-fold smaller as compared to the cLC494. Available gas pressures range between 2 and 50 psi, whereas only from 1 to 3.5 psi in a commercial instrument. Most importantly, only the current system permits reproducible injection after every chemistry cycle of 67% of the end-products in a 5- μ L volume, as compared to 55% in 50 μ L for the cLC494, thereby wholly satisfying a critical requirement for the practical use of any 300-micron ID LC-columns.

Automation control software was written using Labview software on a Macintosh platform (details on software design package, digital control interface and computer are given in '**Materials and Methods**'), and sequencer cycles (about 50-70 min in length; see **Table 2**) developed, allowing fully automated operation of the instrument **100**. In addition to required temperatures, pressure regulation and gas flows, the program controls all rotary plus solenoid valves simultaneously and in a fully coordinated fashion; and allows to string together an essentially unlimited number of "functions" to create cycles of chemistry ("cycles"); which, in turn, can be looped together (i.e. repeated) 20 or more times. Any function itself consists of a particular sequence of unique valve position combinations ("events"), as illustrated in **Figure 2** for function #29, '*deliver liquid TFA to cartridge*', a seven-event function. Here, sequentially, the reaction valves are flushed with argon, TFA metered and delivered to the cartridge, valves flushed, reaction selector valve rinsed with EtAc, flushed with argon and

valves turned into 'safe' position (i.e. central channel in line with plugged port). Operational status can be monitored at all times through the use of a 'virtual instrument' graphical display that emulates actual instrument operation.

Operation and Fine Tuning

The prototype instrument **100** was fully tested for automated operation, followed by optimization of flows and chemistries, and then interfaced with a capillary HPLC system. To this end, precise optimizations of ATZ-extraction/delivery to flask and PTH-aa transfer from flask **101** to injector valve **64** were carried out, largely empirically, with the objective of reducing extraction and transfer volumes. Additional issues addressed were conversion chemistry (conditions of initial aqueous ATZ to PTC conversion, and TFA concentration for generating PTH-aa's), and dissolution (volume; %MeCN/%TFA) and volume reduction (forced argon evaporation) of the PTH-aa's. As a result, the system enables near quantitative LC-injection of volumes in the 5-10 μ L range, making it fully compatible with capillary-LC.

This is most convincingly illustrated in **Figure 3**, where an LC separation of 25 femtomoles of PTH-amino acid standard **15** is shown. Note, that it concerns here 25 femtomoles of standard delivered into the conversion flask, dried down, redissolved, volume reduced by forced evaporation, and injected for analysis. We measured the final volume in the flask to be about 7-8 μ L. Of that final volume, 5 μ L (67%) is then normally injected, representing 17 femtomoles of standard mixture, assuming that no losses have occurred in the process.

Protein Sequencing

Following optimization of the individual chemistry and analysis modules, the integrated system was tested for micro-sequencing performance using, first, 1-pmole and then

400-fmole amounts of non-covalently attached protein standards. Changes were introduced to redress any observed problems until initial yields (i.e. recovery of the first amino acid) were consistently over 50%, repetitive yields (i.e. efficiency of the Edman degradation chemistry) in excess of 90%, and the predominant chemical background (i.e. DPTU) below 1 picomole per cycle. Several thousand cycles have since been completed without any major problems.

An example of the PSMSK cyclic sequencing chemistry, using 100 fmoles of beta-lactoglobulin as starting material, is shown in **Figure 4**. Although we recognize that the lower limits of sensitivity are being approached in this instance, the sequence can actually be called quite well by those skilled in the art, with the exception perhaps for Thr and Lys, residues known to recover at substantially lower yields during Edman degradation.¹⁷ The initial yield was well over 70%, even after background subtraction; repetitive yield, as calculated from the background subtracted yields of PTH-Val in cycles 3 (74 fmoles) and 15 (21 fmoles), was 90.0%. By all means, this would be considered an acceptable result for any sequencing experiment in the event when a system's, any system's, lower limits of sensitivity have been reached. The difference being that in the example presented here, the amount of starting material was about eight to ten times less than hitherto reported for some of the more challenging high-sensitivity experiments using state-of-the-art commercial instruments.^{19,23,24} Our result exceeds manufacturer performance specifications (i.e. 15 residues of 2 pmoles beta-lactoglobulin, with 92% repetitive yield)¹⁹ of an Applied Biosystems cLC494 instrument by at least an order of magnitude.

The PSMSK instrument represents, therefore, a new phase in operative, Edman-chemical protein sequencing, both in terms of miniaturization and sensitivity. To put this advancement in some historical context, a brief overview of the Edman chemistry and its automation during the second half of the 20th century is given in **Table 3**.

Practical Considerations

To function as a truly practical and reliable device for elaborate, constantly repeated cycles of chemistries and analyses, be it process- or micro-scale, certain standard criteria of versatility and routine operation needed to be satisfied. Among PSMSK's most desirable operational features, we will briefly consider its durability and its capacity to utilize large numbers of reagents / solvents in an essentially unlimited sequence of reactions, extractions, drying steps, dissolutions, and transfers.

In the application-validated protein sequencer embodiment described above, ten different 'liquids' 11-16, 141-144 were transferred through diverse combinations of flow-paths and vessels, by using changing gas pressures and valve position combinations, to carry out 86 discrete steps (61 in the reaction cycle; 25 in the conversion cycle -- **Table 1**), totaling 227 'events' per cycle. While permitting a large degree of flexibility already, our automated microfluidics system could easily be upgraded to deliver dozens more of chemicals, through increasingly elaborate flow paths, by adding a few more multi-port selection valves. Such additions would be readily supported by the Labview-based instrument control software:

In cases of frequent usage, it is conceivable that dozens of 'cycles', hundreds of 'steps' and several thousand 'events' would be completed per day, totaling tens of thousands of valve rotations per week, and raising legitimate concerns about abrasion of rotary seals and resultant blockage or leakage. However, this is not what we have observed after more than a year of regular usage, comprising hundreds of thousands of discrete valve movements. Aside from offering some of the lowest internal dead volumes available, 'Valco Cheminert' microbore-type rotary valves (see '**Methods and Materials**') have originally been designed to withstand pressures of up to 3,500 psi and could therefore be slightly loosened (e.g. factory calibrated to

200 psi) to accommodate a maximum pressure of 50 psi in our instrument, thereby extending life expectancies.

Finally, since the instrument is plumbed with capillary tubing, we took extra precautions to prevent salt crystal formation by separating acid and base in the microfluidics system at all times. This was most easily accomplished by isolating TFA 13 on a dedicated valve 66, in conjunction with proper rinsing of the exposed flow paths with EtAc 142 and flushing with argon from three different ports (see Fig. 2). As a result, no particulate-relating problems have ever been observed.

CONCLUSIONS

Detailed biochemical analysis of all molecular communications in a cell will require better tools for microchemical quantitation and identification. We sought to assemble and optimize a microfluidics-based instrument for automation of serial chemical and enzymatic reactions on the smallest possible scale, while maintaining the capacity to analyze a large portion of the end-products. Here we describe a miniaturized, fully integrated and automated system, consisting of multiple rotary valves, reaction and collector modules, all connected by capillary lines, that can deliver about 70% of the end-products, in a 5- μ L volume, to any analytical device with low flow-injection or -infusion (e.g. cLC or NanoESI-MS). A near total control of flow path combinations and directions, temperatures and gas pressures enables precise execution of complex biochemical laboratory procedures. Instrument performance was clearly demonstrated by partially sequencing 100 femtomoles of an intact protein using classical Edman chemistry in combination with capillary-bore liquid chromatographic identification. To our knowledge, this is the smallest amount of protein ever reported to be successfully analyzed in this way. Near quantitative transfer of reaction end-products from a

pressurized micro-vial, in a minimal volume and at exceedingly low flows, could make any biochemical process fully compatible with NanoESI MS or MS/MS analysis;^{31,32} for instance, protein digestion in combination with chemical modification(s).

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Table 1. Comparison of automated protein sequencing instruments: Applied Biosystems 'cLC494' and 'PSMSK'.

RXN, reaction; CNV, conversion; Evalve, cartridge reagent/solvent block or valve; other abbreviations, see key in **Fig. 1** or Glossary. R1 corresponds to **11** in Fig. 1C, R3 to **13**, R4 to **14**, S1 to **144d**, S123 to **each** of **144a**, **144b**, and **144c**.

"Flask" in the table refers to the conversion vessel **101**.

Tables 2A and 2B. Model 'PSMSK' reaction and conversion cycles.

RXN, reaction; CNV, conversion; NLE-Std, PTH-norleucine (30 fmol/ μ L acetonitrile); reaction temperature, 48°C; conversion temperature, 64°C. Total reaction cycle time, ~68 min; total conversion cycle time, ~45 min; further details, see **Fig. 1** and elsewhere in the application.

Table 3. Brief history of Edman protein sequencing.

TABLE 1**A. Wetted Surfaces:**

<u>System</u>	<u>RXN Disc</u>			<u>RXN Block</u>
	<u>Diameter</u>	<u>Surface</u>	<u>Volume</u>	<u>Cavity Volume</u>
	<i>mm</i>	<i>mm²</i>	<i>μL</i>	<i>μL</i>
cLC494	6	28	8	32
PSMSK	3	7	2	4

<u>System</u>	<u>Flow Path Volume</u>			<u>Injector</u>
	<u>Evalve-Flask</u>	<u>Flask-Injector</u>	<u>Total</u>	<u>Loop Volume</u>
	<i>μL</i>	<i>μL</i>	<i>μL</i>	<i>μL</i>
cLC494	248	130	378	50
PSMSK	11	1.5	12.5	5

TABLE 1 (CONTINUED)**B. Reagents/Solvents Delivery:**

<u>System</u>	<u>Disc</u> <i>mm</i>	<u>Polybrene</u> <i>mg</i>	<u>RXN</u>		<u>WASH (S123)</u>	
			<u>R1</u>	<u>R3</u>	<u>Volume</u>	<u>Vol./Area</u>
			μL	μL	μL	$\mu\text{L}/\text{mm}^2$
cLC494	6	0.75	8	32	920	33
PSMSK	3	0.18	2	4	270	39

<u>System</u>	<u>ATZ Transfer</u>		<u>CNV</u>			<u>Injection</u>
	<u>Volume</u>	<u>Vol.Area</u>	<u>S4(preATZ)</u>	<u>R4</u>	<u>S4(PTH)</u>	<u>Percentage</u>
	μL	$\mu\text{L}/\text{mm}^2$	μL	μL	μL	%
cLC494	291	10.4	60	10	91	55
PSMSK	76	10.9	7.5	7.5	7.5	67

TABLE 2AReaction Cycle - #27 Glass Fiber

<u>Step</u>	<u>Description</u>	<u>Time Elapsed (sec)</u>	
		<u>Core</u>	<u>Step</u>
1	Deliver TMA to Cartridge	20	68
2	Deliver PITS to Cartridge	10	35
3	Pause	20	20
4	Deliver TMA to Cartridge	400	448
5	Deliver PITS to Cartridge	10	35
6	Pause	20	20
7	Deliver TMA to Cartridge	400	448
8	Deliver PITS to Cartridge	10	35
9	Pause	20	20
10	Deliver TMA to Cartridge	400	488
11	AR4 to Cartridge	240	248
12	Deliver Heptane to Cartridge	60	83
13	Pause	10	10
14	AR4 to Cartridge	10	18
15	Deliver Heptane to Cartridge	60	83
16	Pause	10	10
17	AR4 to Cartridge	10	18
18	Deliver BuCl to Cartridge	60	83
19	Pause	10	10
20	AR4 to Cartridge	10	18
21	Deliver EtAc to Cartridge	60	83
22	Pause	10	10
23	AR4 to Cartridge	10	18
24	Deliver EtAc to Cartridge	60	83
25	Pause	10	10
26	AR4 to Cartridge	10	18
27	Deliver EtAc to Cartridge	60	83
28	Pause	10	10
29	AR4 to Cartridge	150	158
30	Deliver Liquid TFA to Cartridge	10	104
31	Load RXN Loop with EtAc	15	23
32	Clear RXN Loop to Waste with AR5		10
33	Load RXN Loop with EtAc	15	23
34	Clear RXN Loop to Waste with AR5		40
35	AR4 to Cartridge	20	248
36	Prep Transfer	10	10
37	Deliver Heptane to Midpoint	12	32

38	Transfer Pause		15	15
39	Transfer with BuCl Part 1	15	15	
40	Transfer Pause		15	15
41	Transfer with AR4	45	55	
42	Transfer Argon Pulse	40	40	
43	Transfer with BuCl Part 1	15	15	
44	Transfer Pause		10	10
45	Transfer with AR4	30	40	
46	Transfer Argon Pulse	30	30	
47	Transfer with BuCl Part 1	15	15	
48	Transfer Pause		10	10
49	Transfer with AR4	30	40	
50	Transfer Argon Pulse	30	30	
51	Transfer with BuCl Part 1	15	15	
52	Transfer Pause		10	10
53	Transfer with AR4	30	40	
54	Transfer Argon Pulse	30	30	
55	End Transfer	1	1	
56	Deliver BuCl to Cartridge	30	53	
57	AR4 to Cartridge	120	128	
58	Load RXN loop with BuCl	60	68	
59	Load RXN loop with Heptane	60	68	
60	Load RXN loop with EtAc	60	68	
61	AR4 to Cartridge	120	128	

End of Table 2A

TABLE 2B
Conversion Cycle - #28 Low Pressure Conversion

<u>Step</u>	<u>Description</u>	<u>Time Elapsed (sec)</u>	
		<u>Core</u>	<u>Step</u>
1	Empty Flask to Injector	60	68
2	Rinse Vent/Flask with 25% TFA	45	72
3	Rinse Vent/Flask with 25% TFA	45	72
4	Dry Flask with AR6	60	68
5	Empty Flask to Waste with AR6	400	408
6	Rinse Vent/Flask with 10%MeCN	30	52
7	Rinse Vent/Flask with 10%MeCN	30	52
8	Dry Flask with AR6	60	68
9	Empty Flask to Waste with AR6	300	308
10	Dry Flask with AR5	60	68
11	Load CNV Loop with NLE-Std	10	16
12	Deliver CNV Loop to Flask with AR5	80	80
13	Load CNV Loop with 25%TFA	15	21
14	Deliver CNV Loop to Flask with AR5	20	20
15	Ready to Receive	1	1
16	Deliver CNV Loop to Flask with AR4	1000	1000
17	Load Injector Loop with AR5		240 248
18	Load CNV Loop with 10%MeCN	10	16
19	Deliver CNV Loop to Flask with AR5	20	20
20	Pause	10	10
21	Deliver CNV Loop to Flask with AR5	20	20
22	Pause	10	10
23	Load Injector Loop with AR5		15 23
24	Relay 1 On: Start HPLC	2	2
25	Relay 1 Off	2	2

TABLE 3**Brief History of Edman Protein Sequencing**

<u>YEAR</u>		<u>SCALE</u>	<u>REFERENCES</u>
1950	Edman Chemistry	1 mmol	9,10
1955-56	Practical Manual Sequencing	1 mmol	33
1967	Spinning-Cup Sequencer	200 nmol	11
1968-77	Various Improvements	10-100 nmol	13,34
1978	Picomole HPLC analysis	0.5-5 nmol	14
1981	Gas-Phase Sequencer	20-100 pmol	15
1986	'On-line' HPLC	5-10 pmol	
1987-94	Various Optimizations	2-5 pmol	16,17,18
1996	Applied Biosystems cLC	1 pmol	19
2000	Microfluidics (PSMSK)	100-200 fmol	this patent application

What is claimed is:

1. A system for carrying out one or more chemical reactions, said system comprising a rotary selector valve and a rotary switching valve, each valve under the control of a computer, the internal volumes of the selector and switching valves each being 1.5 μL or less.

2. A system of Claim 1 wherein the rotary selector valve comprises, in a first unit, a central common port and a circular array of a plurality of peripheral ports such that the main axis of the common port is the same as the main axis through the circular array, and wherein said valve comprises, on a second unit, a radial connector channel, the first and second units being juxtaposed such that there is a single continuous selector channel formed by the common port, the connector channel, and a peripheral port, the selection of the peripheral port under control of the computer, the internal volume of the continuous selector channel being the internal volume of the valve, and

wherein the rotary switching valve comprises a circular array of three or more peripheral ports in a first unit and a connector channel in a second unit, the two units being juxtaposed such that there is a continuous switching channel formed by a first peripheral port, the connector channel, and a second peripheral port, the selection of two peripheral ports being under the control of the computer, the internal volume of the continuous switching channel being the internal volume of the valve.

3. A system of Claim 2 wherein the rotary selector valve is connected by its central common port to a first peripheral port of the rotary switching valve, the connection being accomplished by a conduit means having an internal volume between 0.5 μL and 10 μL .

4. A system of Claim 3 wherein the total internal volume of the selector and switching valves are each in the range 0.04 μL to 1.5 μL .

5. A system of Claims 4 wherein the total internal volume of the selector and switching valves are in the range 0.1 μL to 0.5 μL .

6. A system of Claim 3, said system further comprising :
a reaction vessel connected to the second peripheral port in the continuous switching channel of the switching valve,
a first and second reagent vessel, said vessels each connected to first and second peripheral ports of the selector valve, said vessels each connected to a gas source comprising a gas under pressure.

7. A system of Claim 6, the selector valve thereof being a first selector valve and the switching valve thereof being a first switching valve, said system further comprising:
a second rotary selector valve connected to a second rotary switching valve;
a third rotary switching valve;
a conversion vessel connected to receive output from said third rotary switching valve,
a restraining means for restraining unprocessed polymer in the reaction vessel;
third and fourth reagent vessels connected to provide fluid to said conversion vessel;
wherein said second rotary valve, second switching valve, and third switching valve are each under the control of the computer and each have an internal volume of 1.5 μl or less.

8. A reaction process under the control of a computer said process comprising the steps of:

- 1) Delivering, from a reagent vessel under pressure from a gas source, a volume of a reagent via a delivery line to a reaction vessel wherein the total volume delivered for purposes of a reaction in the reaction vessel is in the range 0.5 uL to 10 uL, and wherein flow through the delivery line is under the control of a computer-controlled rotary switching valve connected to a rotary selector valve.

9. The process of Claim 8, said process further comprising steps (2) and (3):

- 2) Washing said valves and delivery line with a wash-solvent; and
- 3) Repeating step (1) optionally with a reagent, reagent vessel and/or gas source different from that used in step (1).

10. A process of Claim 9, said process further comprising the steps of:

- (4) Placing a polymer in a reaction vessel, said polymer restrained in said vessel;
- (5) Delivering to said reaction vessel, from a first reagent vessel under pressure from a gas source, a volume of a first fluid, said fluid comprising a first reagent, which reagent will bind to a terminal monomer on the polymer;
- (6) Delivering to said reaction vessel, from a second reagent vessel under pressure from a gas source, a volume of a second fluid, which fluid will cause a terminal monomer-reagent moiety to be cleaved from the polymer;
- (7) causing all or part of the fluid in the reaction flask to be transferred under gas pressure via a delivery line to a conversion flask while allowing the polymer, less the terminal monomer, to remain in the reaction vessel;

- (8) Prior or after step (7), delivering from a third reagent vessel under pressure from a gas source a volume of a third fluid such that the combination of said fluid and the fluid transferred in step (7) to the reaction flask will cause the terminal monomer-reagent moiety to be cleaved to create a terminal monomer free of covalently bound reagent;
- (9) Transferring under gas pressure from a gas source said terminal monomer created in step (8) to an analytical device that will identify the nature of the terminal monomer;

wherein each volume delivered in steps (5), (6), (7), (8), is in the range 0.2 uL to 10 uL, and the volume delivered in step (9) is in the range 1 uL to 10 uL.

11. A process of Claim 10 wherein steps (4) - (9) are performed a plurality of times.

12. A process of Claim 8 wherein the process comprises sequentially delivering acid, organic solvent, and base to the reaction vessel.

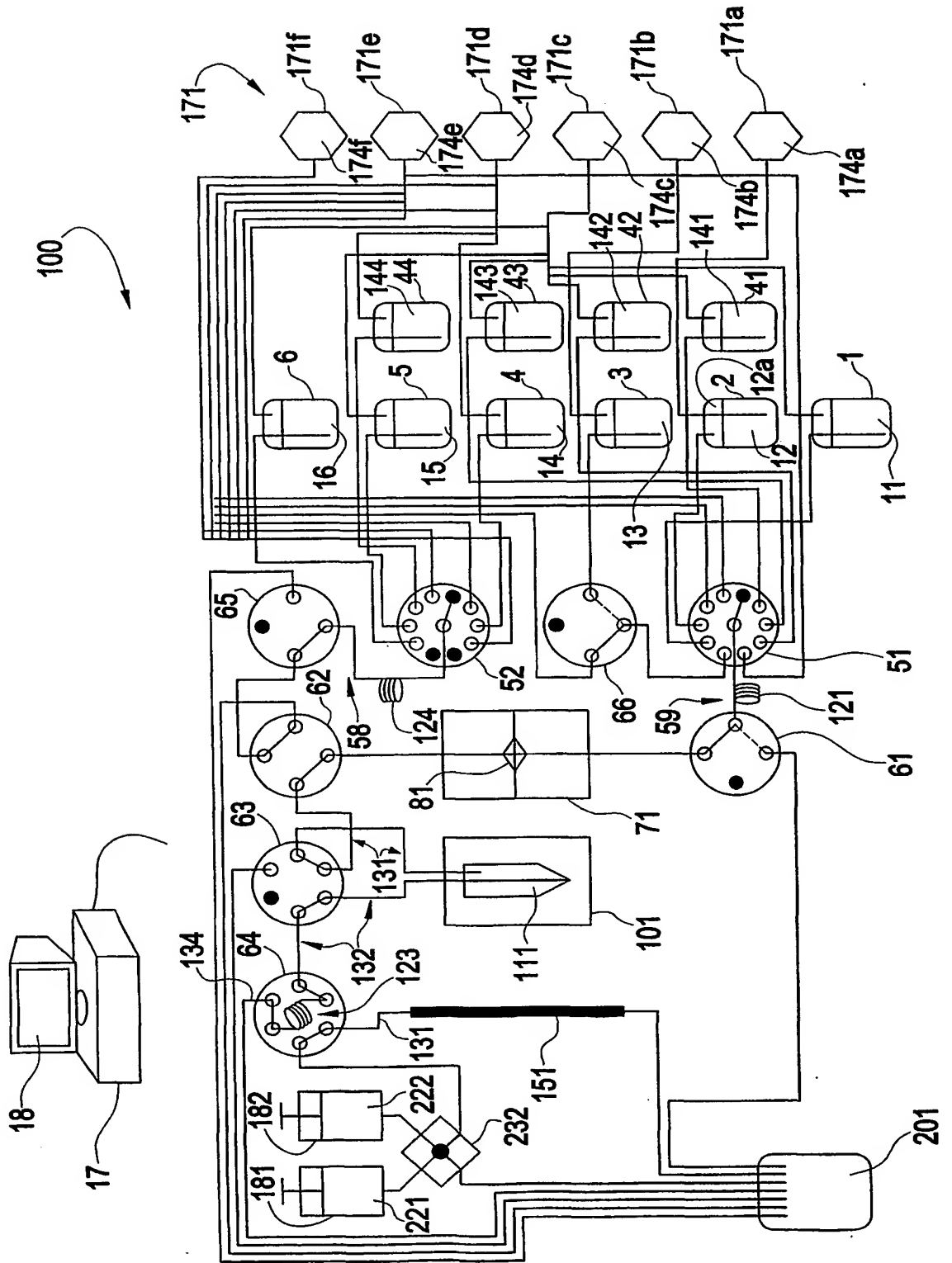
13. A process of Claim 8 wherein the reaction vessel is kept free of air and oxygen.

14. A process of Claim 10 wherein the reaction vessel and the conversion vessel are kept free of air and oxygen.

15. A rotary selector valve cascade in which a plurality of peripheral ports of a first selector valve are each connected to the central common port of one of a plurality of rotary

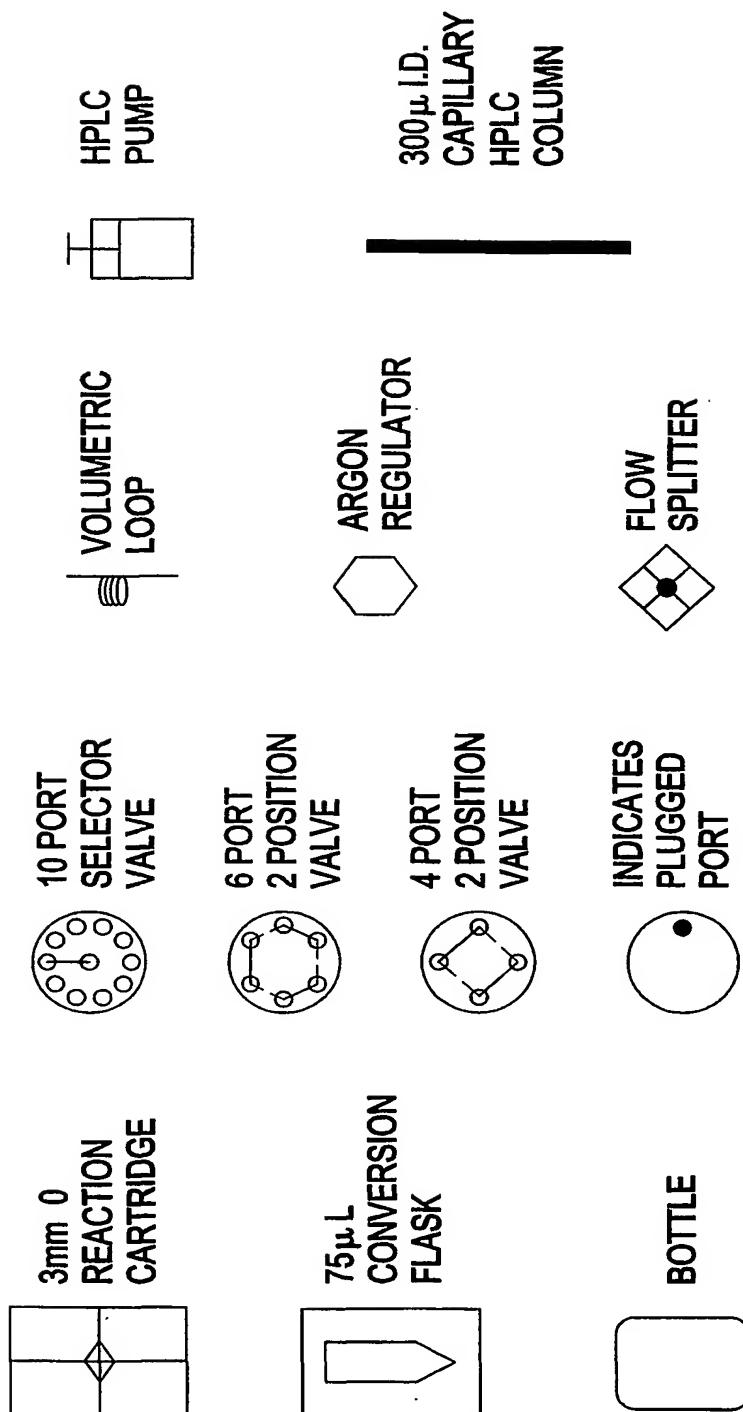
selector valves that each comprise a plurality of peripheral ports, the total internal volume of each valve being 1.5 uL or less.

FIG. 1A



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FIG. 1B



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FIG. 1C

EXAMPLES OF REAGENTS

11	1% PHENYL ISOTHIOCYANATE IN HEPTANE
12	12.5% TRIMETHYLAMINE IN WATER
13	TRIFLUOROACETIC ACID
14	25% TRIFLUOROACETIC ACID IN WATER
15	PTH AMINO ACID STANDARD
16	NORLEUCINE STANDARD

EXAMPLES OF WASH SOLVENTS

141	HEPTANE
142	ETHYL ACETATE
143	BUTYL CHLORIDE
144	10% ACETONITRILE IN WATER
221	WATER WITH TETRAHYDROFURAN
222	ACETONITRILE

EXAMPLES OF GAS SOURCES

174a	2.0 PSI ARGON
174b	2.0 PSI ARGON
174c	4.0 PSI ARGON
174d	12.0 PSI ARGON
174e	24.0 PSI ARGON
174f	50.0 PSI ARGON

FIG. 2B

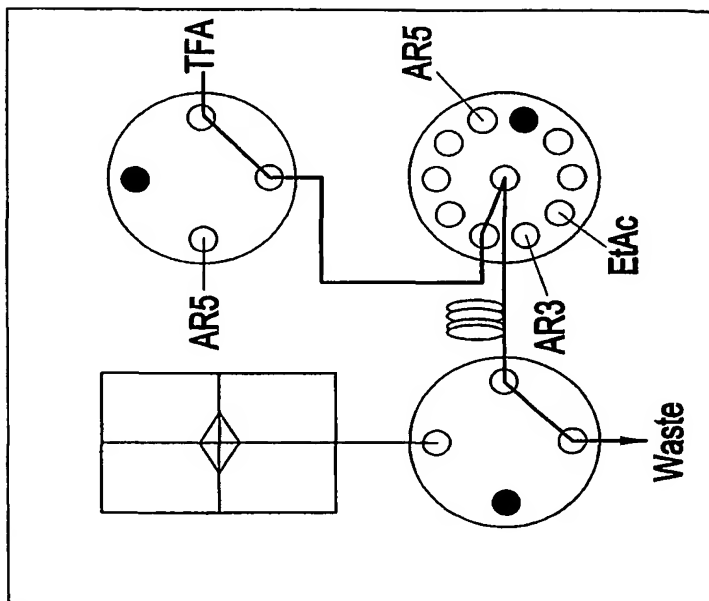


FIG. 2A

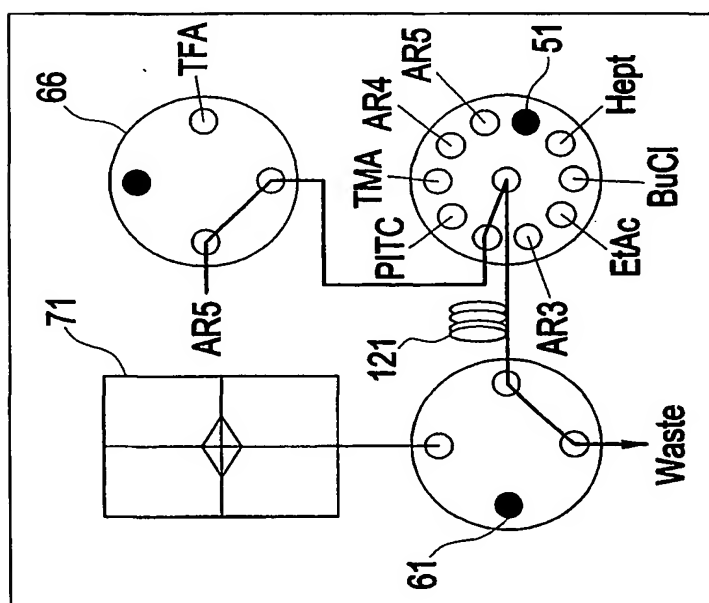


FIG. 2D

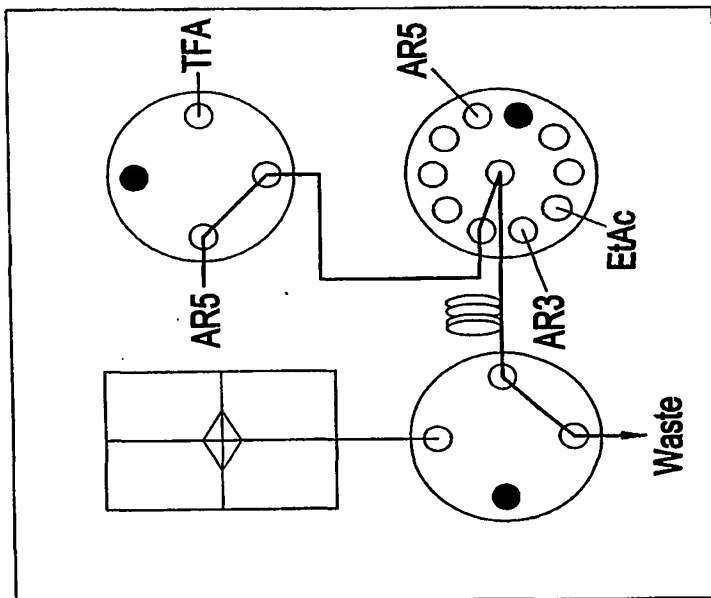
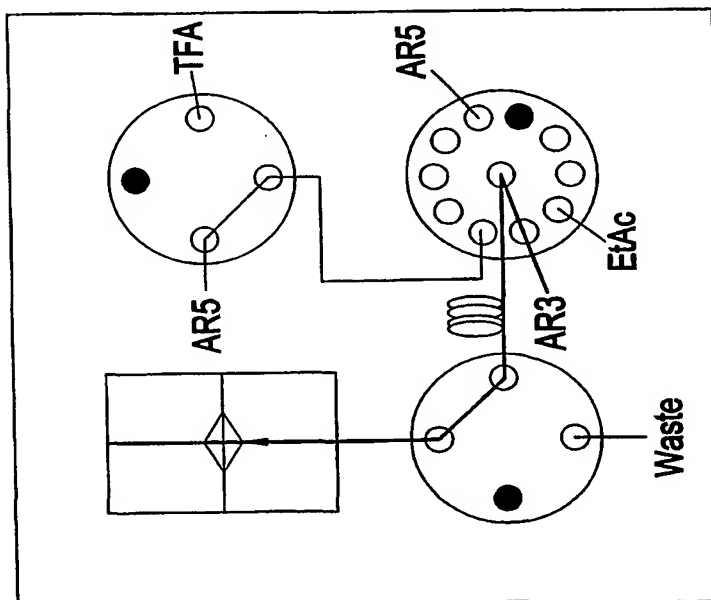


FIG. 2C



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FIG. 2F

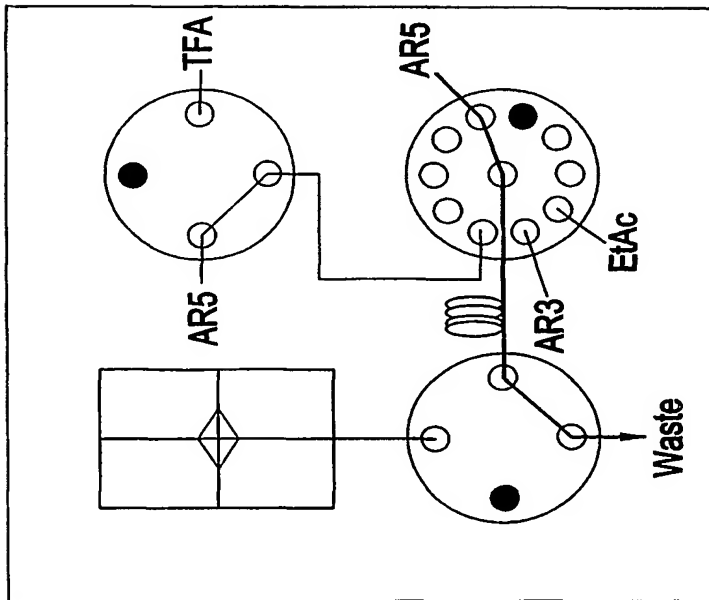
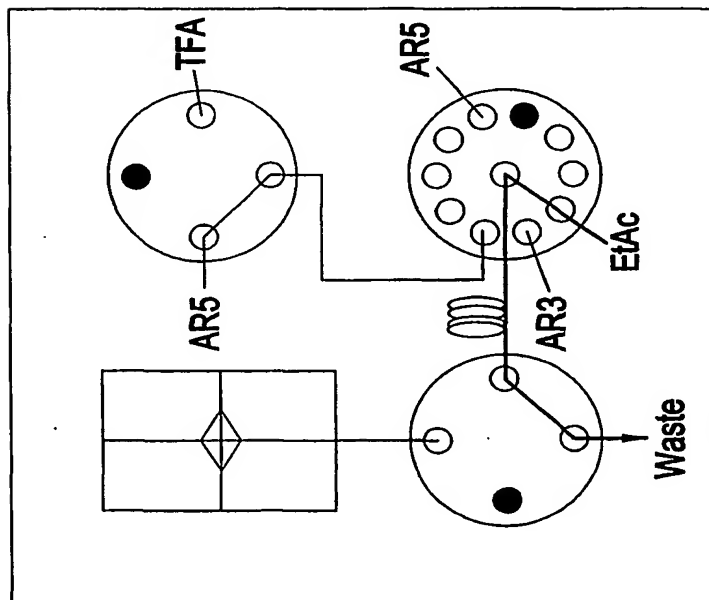
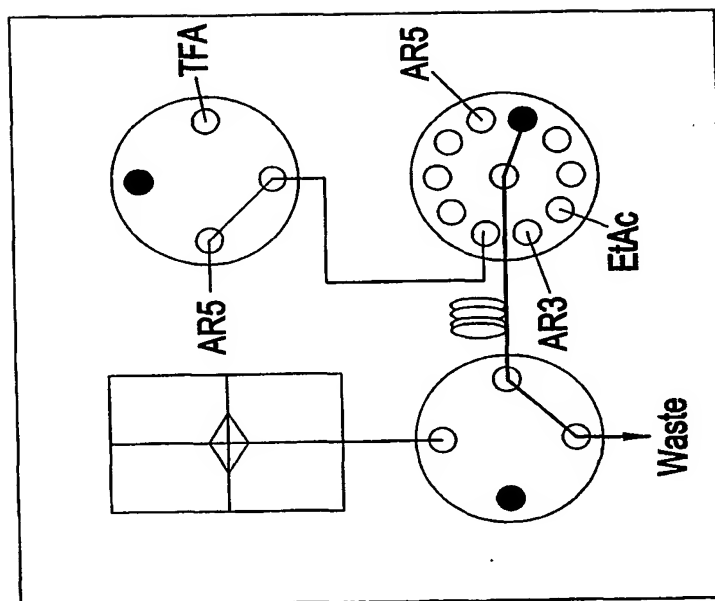


FIG. 2E



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FIG. 2G



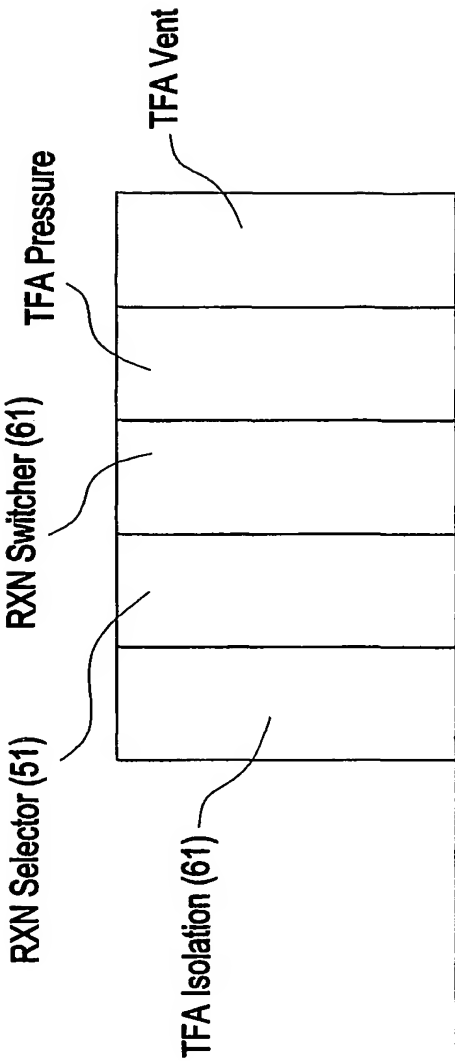
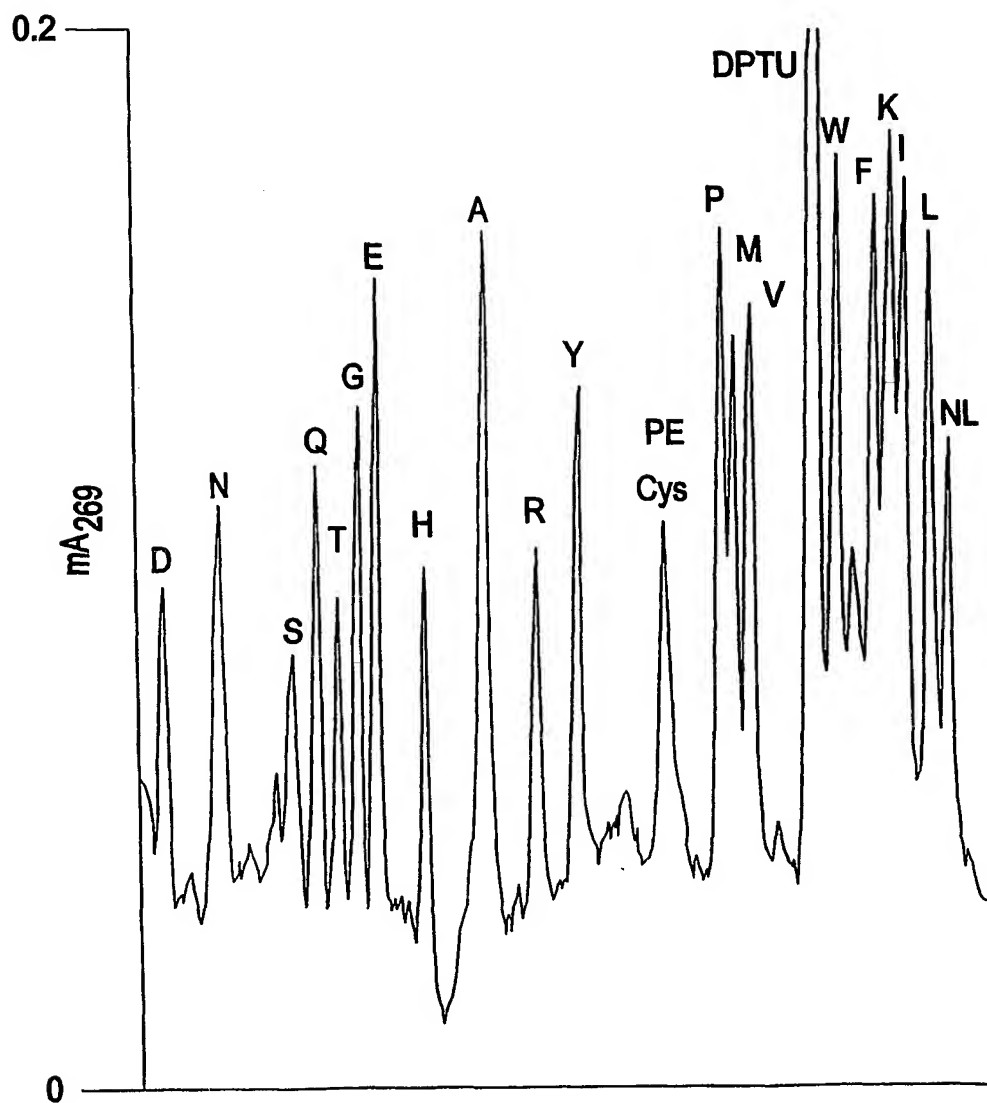


FIG. 2H

EVENT	TIME	DESCRIPTION	VALVE POSITION			
1	10	Preflush RXN Valves with Argon 5	A	6	A	ON
2	20	Load RXN Loop with TFA	B	6	A	ON
3	10	Deliver Loop of TFA to RXN Cartridge	A	7	B	OFF
4	3	Postflush RXN Valves with Argon 5	A	6	A	OFF
5	30	Rinse RXN Selector with EtAc	A	8	A	OFF
6	30	Flush RXN Valves with Argon 5	A	2	A	OFF
7	1	All RXN Valves in Safe Position	A	1	A	OFF

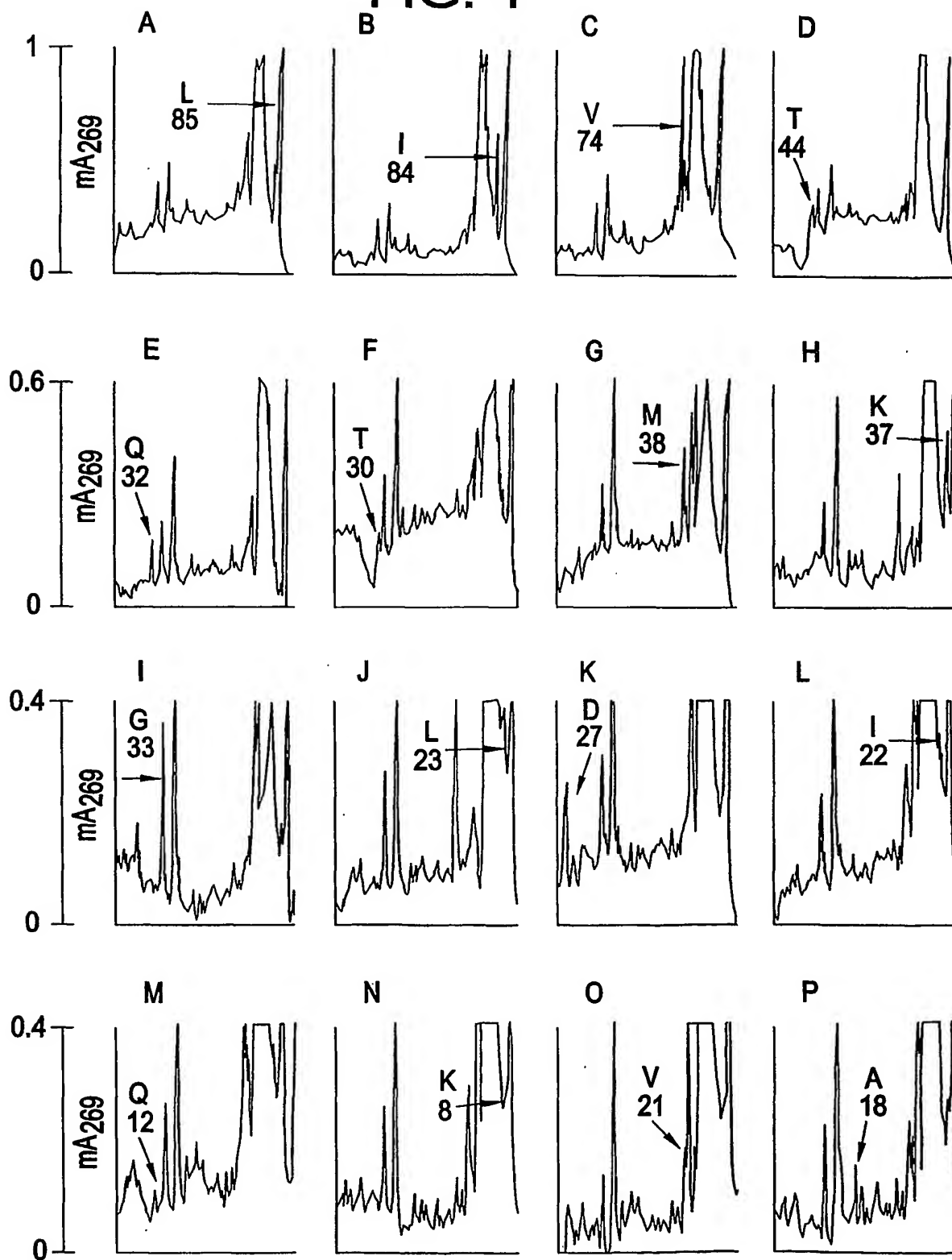
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FIG. 3



SUBSTITUTE SHEET (RULE 26)

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FIG. 4

SUBSTITUTE SHEET (RULE 26)

FIG. 5

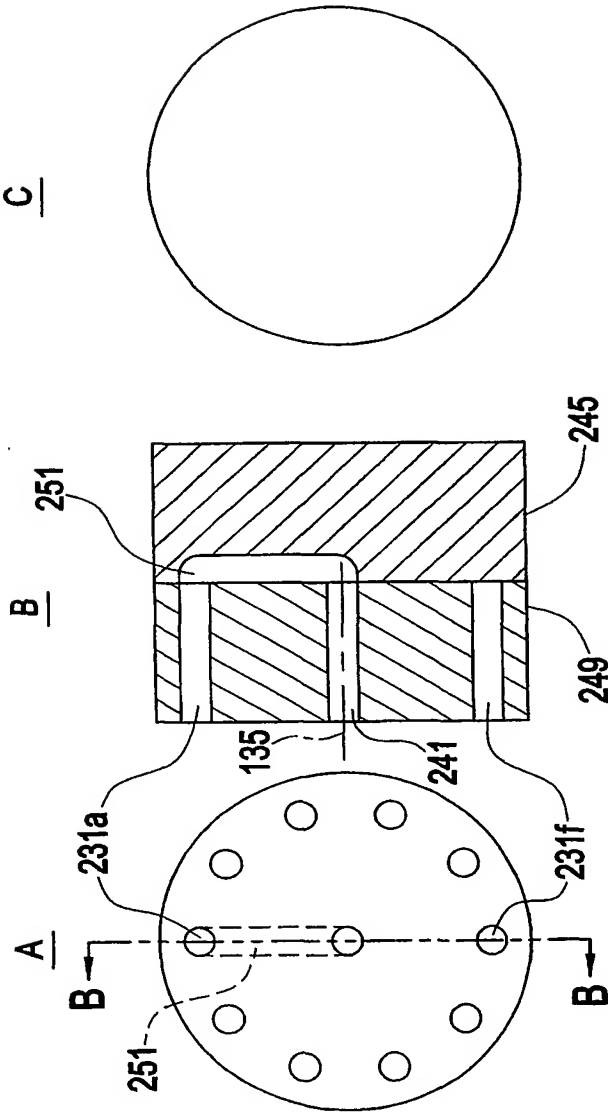
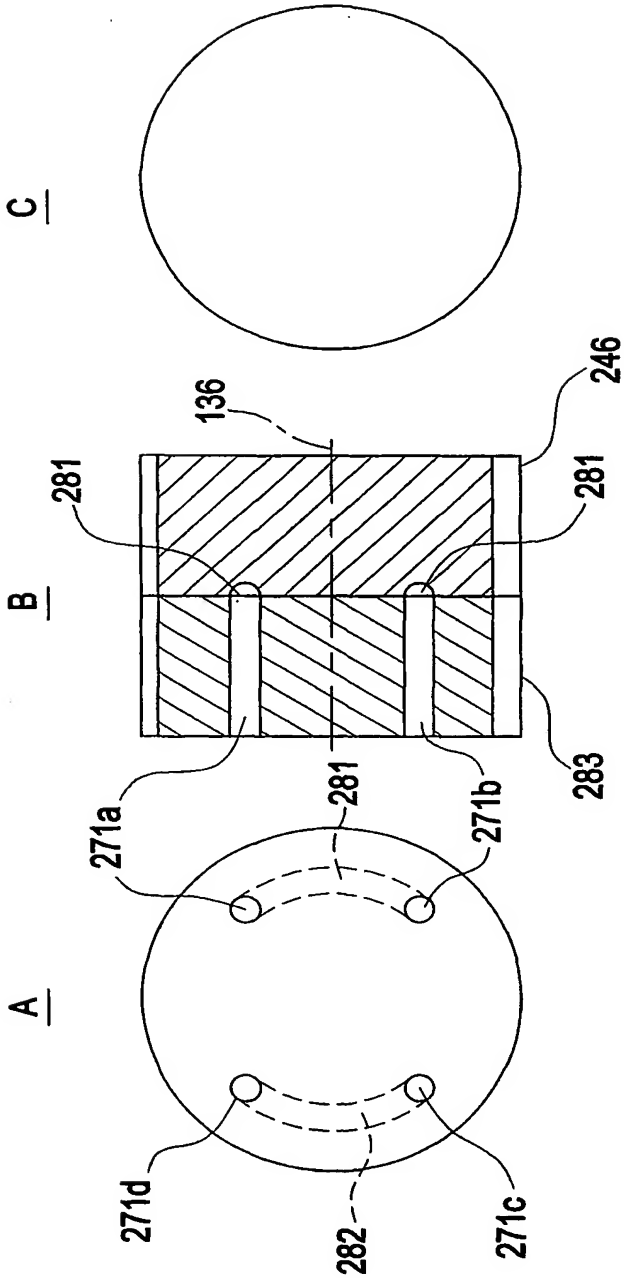
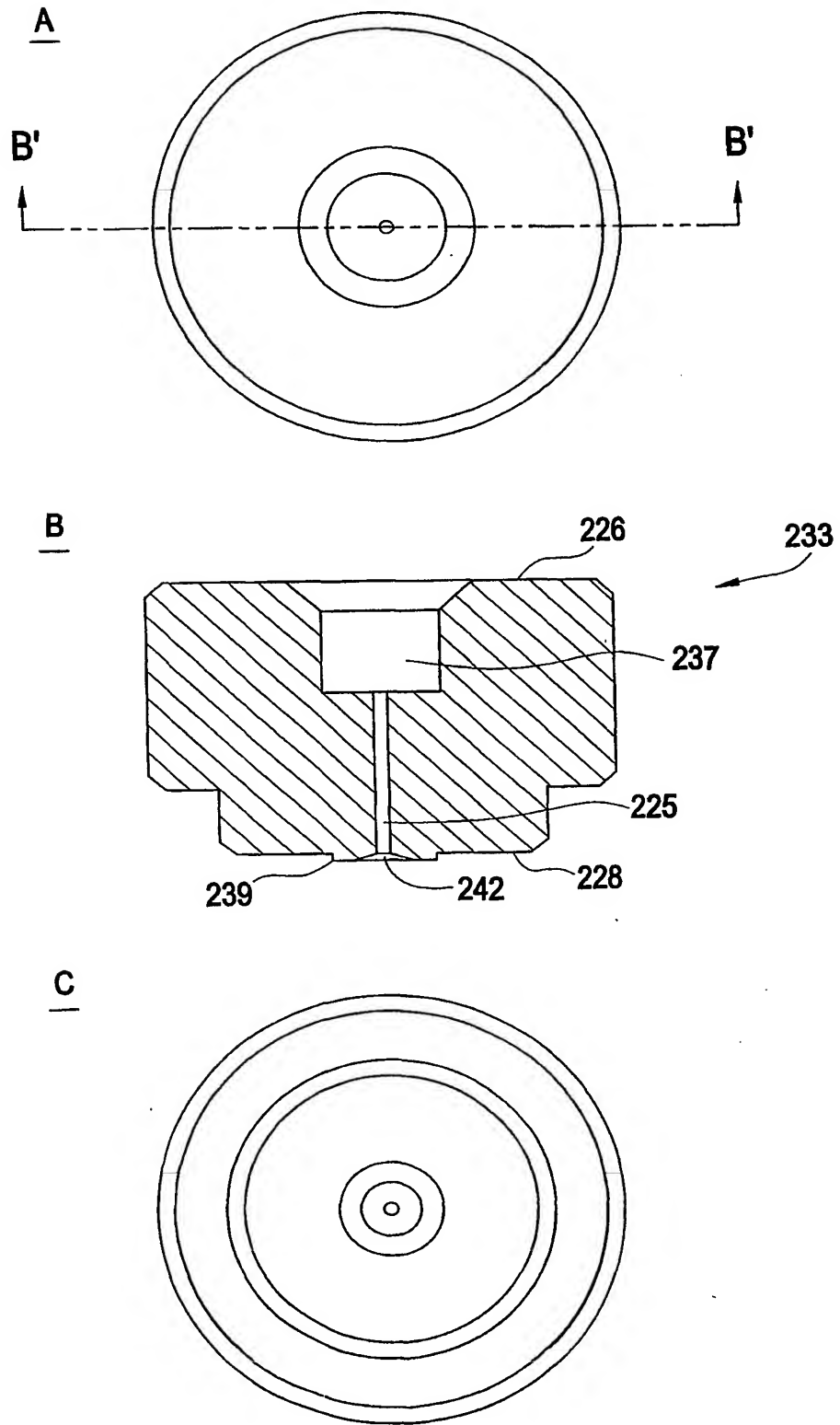


FIG. 6



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FIG. 7



SUBSTITUTE SHEET (RULE 26)

FIG. 8

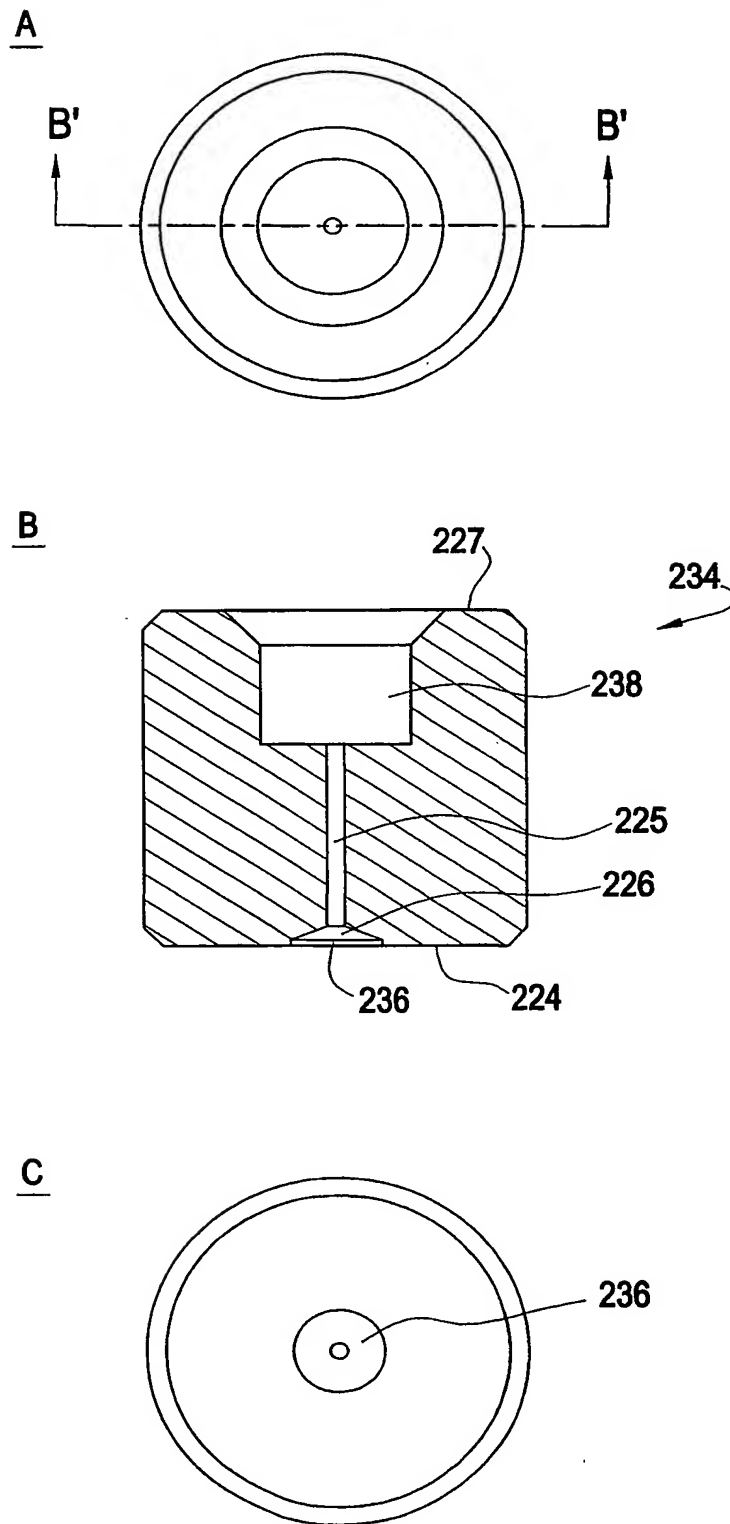


FIG. 9

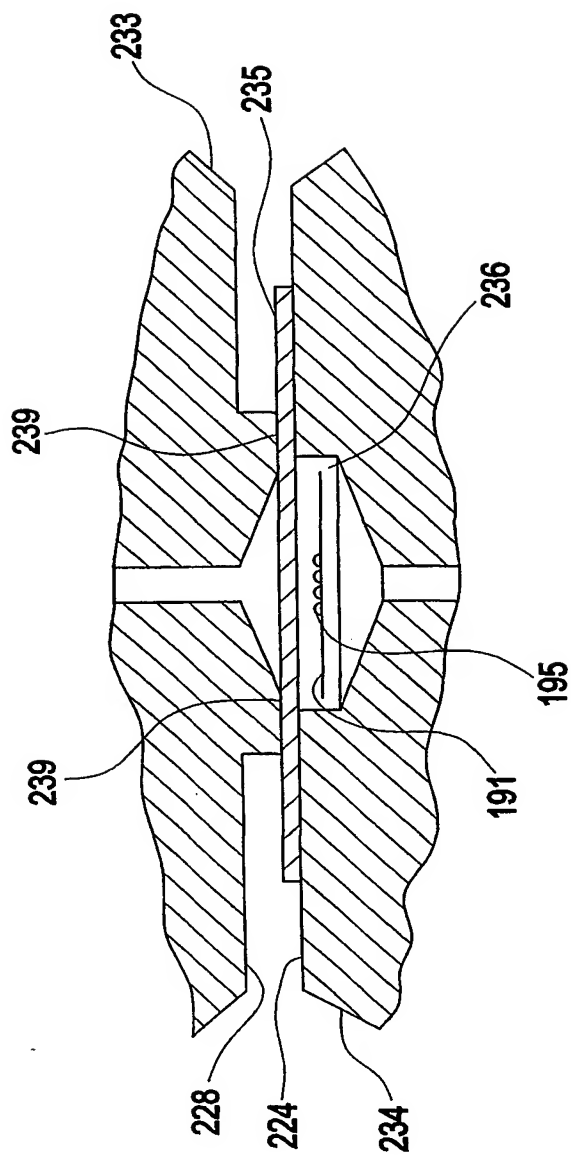


FIG. 10

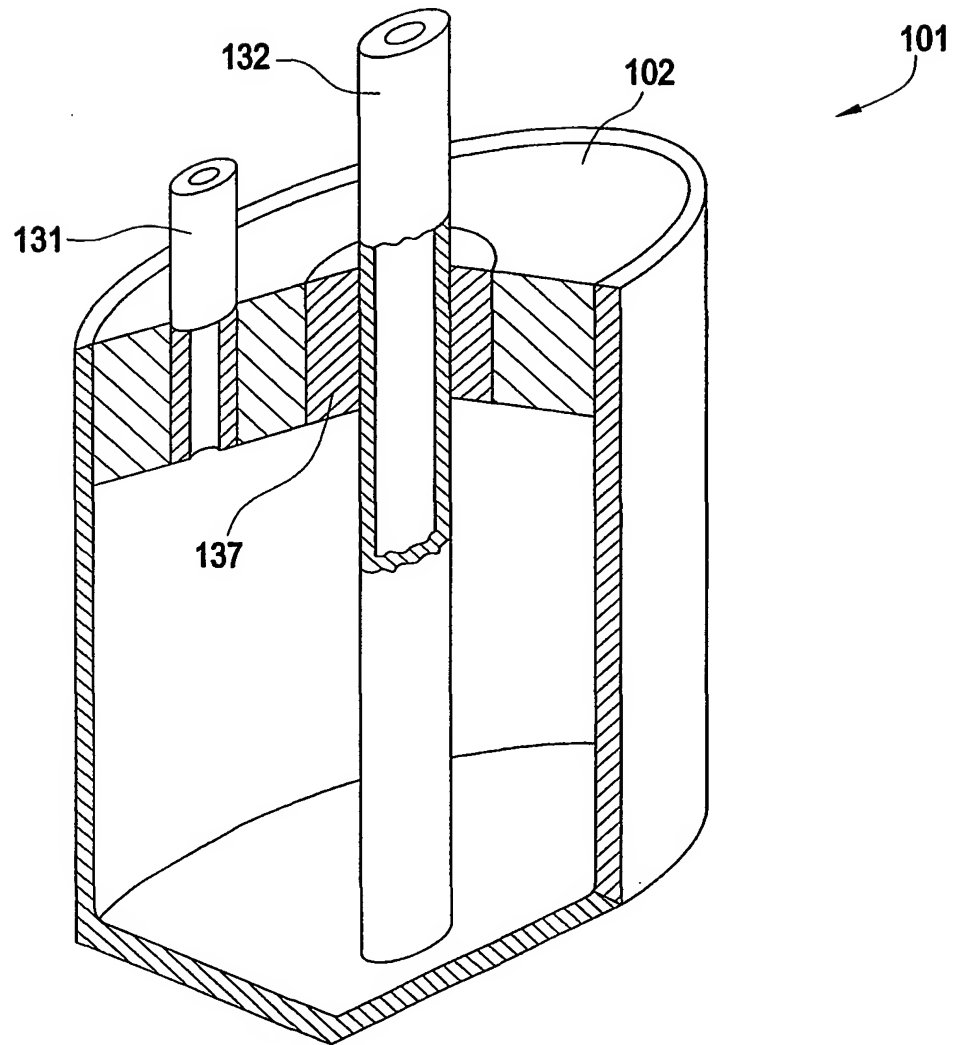


FIG. 11

